

This Page Is Inserted by IFW Operations
and is not a part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

IMAGES ARE BEST AVAILABLE COPY.

**As rescanning documents *will not* correct images,
please do not report the images to the
Image Problem Mailbox.**



1/75

FIG. 1A

2H7scFv-Ig cDNA and predicted amino acid sequence:

HindIII	NcoI	2H7 V _L Leader Peptide→
-----	-----	
1 AAGCTTGCCG CC	ATGGATTTCAGCTTCAGCTTAAT	M D F Q V Q I F S F L L I S A S CAGTGCTTCA

2H7 V_L →
 V I I A R G Q I V L S Q S P A I L S A S
 61 GTCATAATTG CCAGAGGACA AATTGTTCTC TCCCAGTCTC CAGCAATCCT GTCTGCATCT
 P G E K V T M T C R A S S S V S Y M H W
 121 CCAGGGGAGA AGGTACAAAT GACTTGCAGG GCCAGCTCAA GTGTAAGTTA CATGCACTGG

BamHI

 Y Q Q K P G S S P K P W I Y A P S N L A
 181 TACCAGCAGA AGCCAGGATC CTCCCCAAA CCCTGGATTG ATGCCCATC CAACCTGGCT
 S G V P A R F S G S G S G T S Y S L T I
 241 TCTGGAGTCC CTGCTCGCTT CAGTGGCAGT GGGTCTGGGA CCTCTTACTC TCTCACAAATC
 S R V E A E D A A T Y Y C Q Q W S F N P
 301 AGCAGAGTGG AGGCTGAAGA TGCTGCCACT TATTACTGCC AGCAGTGGAG TTTAACCCA

(Gly₄Ser)₃ Linker
 P T F G A G T K L E L K G G G G S G G G
 361 CCCACGTTCG GTGCTGGAC CAAGCTGGAG CTGAAAGGTG CGGGTGGCTC GGGCGGTGGT

2H7 V_H →
 G S G G G G S S Q A Y L Q Q S G A E L V
 421 GGATCTGGAG GAGGTGGAG CTCTCAGGCT TATCTACAGC AGTCTGGGC TGAGCTGGT
 R P G A S V K M S C K A S G Y T F T S Y
 481 AGGCCTGGGG CCTCAGTGAA GATGTCTGC AAGGCTTCTG GCTACACATT TACCAAGTTAC
 N M H W V K Q T P R Q G L E W I G A I Y
 541 AATATGCACT GGGTAAAGCA GACACCTAGA CAGGGCCTGG AATGGATTGG AGCTATTAT
 P G N G D T S Y N Q K F K G K A T L T V
 601 CCAGGAAATG GTGATACTTC CTACAATCAG AAGTTCAAGG GCAAGGCCAC ACTGACTGTA
 D K S S S T A Y M Q L S S L T S E D S A
 661 GACAAATCCT CCAGCACAGC CTACATGCAG CTCAGCAGCC TGACATCTGA AGACTCTGCG
 V Y F C A R V V Y Y S N S Y W Y F D V W
 721 GTCTATTCT GTGCAAGAGT GGTGTACTAT AGTAACTCTT ACTGGTACTT CGATGTCTGG

FIG. 1B

BclI
-----human IgG1 Fc domain →

781 G T G T T V T V S D Q E P K S C D K T H
 GGCACAGGG A CCACGGTCAC CGTCTCTGAT CAGGAGCCCA AATCTTGTGA CAAAACCTCAC

841 T C P P C P A P E L L G G P S V F L F P
 ACATGCCAC CGTGCCCAGC ACCTGAACTC CTGGGGGAC CGTCAGTCTT CCTCTTCCCC

901 P K P K D T L M I S R T P E V T C V V V
 CCAAAACCCA AGGACACCCT CATGATCTCC CGGACCCCTG AGGTACACATG CGTGGTGGT

961 D V S H E D P E V K F N W Y V D G V E V
 GACGTGAGCC ACGAAGACCC TGAGGTCAAG TTCAACTGGT ACGTGGACGG CGTGGAGGTG

1021 H N A K T K P R E E Q Y N S T Y R V V S
 CATAATGCCA AGACAAAGCC GCAGGAGGAG CAGTACAACA GCACGTACCG TGTGGTCAGC

1081 V L T V L H Q D W L N G K E Y K C K V S
 GTCTCACCG TCCTGCACCA GGACTGGCTG AATGGCAAGG AGTACAAGTG CAAGGTCTCC

1141 N K A L P A P I E K T I S K A K G Q P R
 AACAAAGCCC TCCCAGCCCC CATCGAGAAA ACAATCTCCA AAGCCAAAGG GCAGCCCCGA

1201 E P Q V Y T L P P S R D E L T K N Q V S
 GAACCACAGG TGTACACCCT GCCCCCATCC CGGGATGAGC TGACCAAGAA CCAGGTCAAGC

1261 L T C L V K G F Y P S D I A V E W E S N
 CTGACCTGCC TGGTCAAAGG CTTCTATCCC AGCGACATCG CCGTGGAGTG GGAGAGCAAT

1321 G Q P E N N Y K T T P P V L D S D G S F
 GGGCAGCCGG AGAACAACTA CAAGACCACG CCTCCCGTGC TGGACTCCGA CGGCTCCTTC

1381 F L Y S K L T V D K S R W Q Q G N V F S
 TTCCTCTACA GCAAGCTCAC CGTGGACAAG AGCAGGTGGC AGCAGGGAA CGTCTTCTCA

1441 C S V M H E A L H N H Y T Q K S L S L S
 TGCTCCGTGA TGCATGAGGC TCTGCACAAC CACTACACGC AGAAGAGCCT CTCCCTGTCT

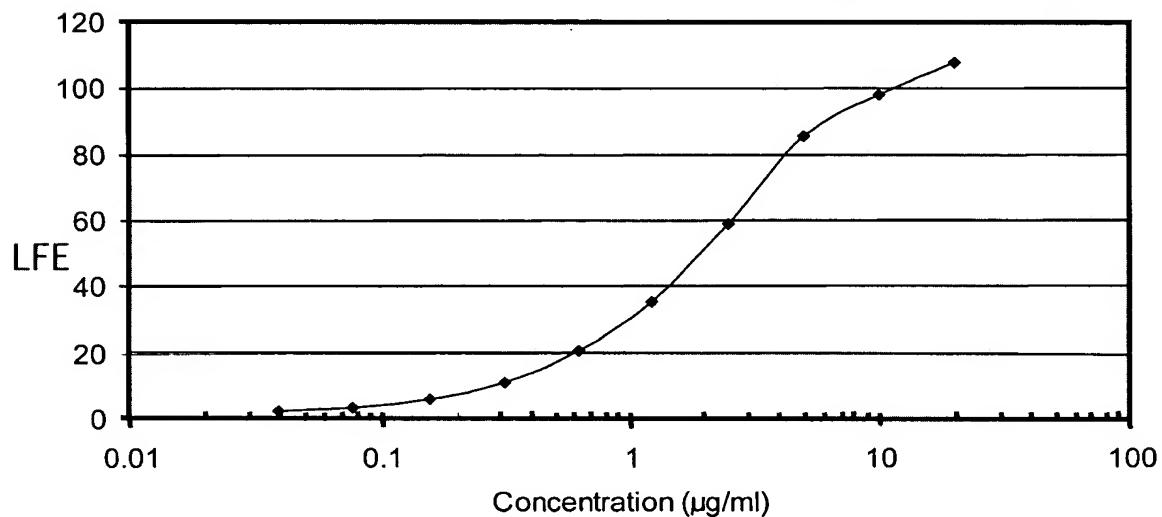
XbaI

1501 P G K * S R
 CCGGGTAAAT GATCTAGA

FIG. 2

Production Levels of 2H7 scFv (SSS-S)H WCH2 WCH3
by Stable CHO Lines

2H7 scFv (SSS-S)H WCH2 WCH3 STANDARD CURVE



Clone	LFE @ 1:50 Estimated Concentration (mg/ml)
D2	26.156
IIIC6	25.755
IVA3	28.661
Spent bulk	29.664

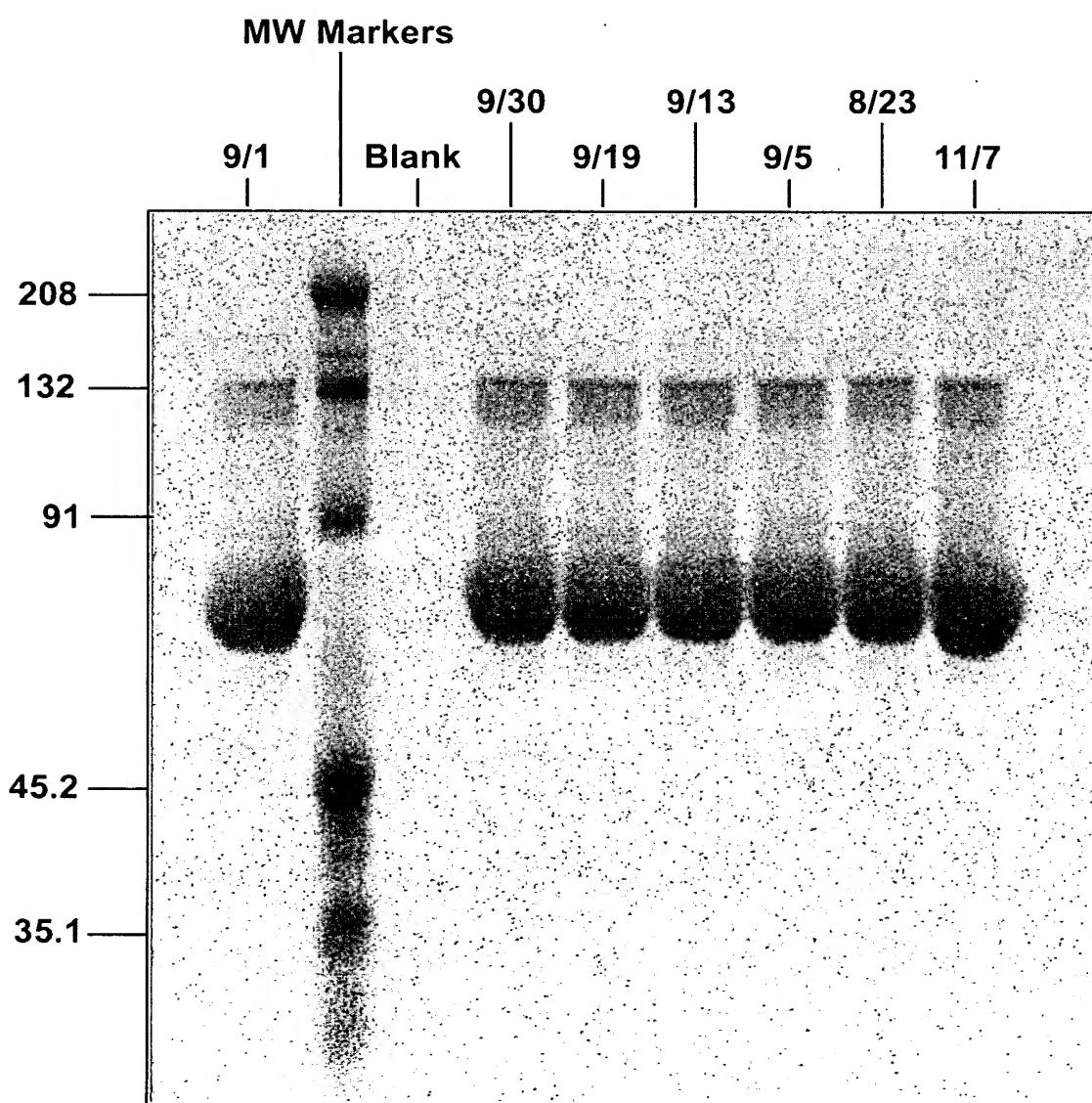
FIG.3**SDS-Page Analysis of
2H7 scFv (SSS-S)H WCH2 WCH3 Protein**

FIG. 4A

**Complement Mediated B Cell Killing After Binding of CD20-targeted
2H7 scFv (SSS-S)H WCH2 WCH3**

2H7scFv-Ig Concentration	RAMOS		BJAB	
	# LIVE CELLS / TOTAL CELLS		# LIVE CELLS / TOTAL CELLS	
20 µg/ml + complement	—	0.16	—	0.07
5 µg/ml + complement	—	0.2	—	N.D.
1.25 µg/ml + complement	—	0.32	—	0.1
Complement alone	—	0.98	—	0.94

*Viability was determined by trypan blue exclusion and is tabulated as the fraction of viable cells out of the total number of cells counted.

**N.D. (not determined).

FIG. 4B

Antibody-dependent cellular cytotoxicity (ADCC) mediated by
2H7 scFv (SSS-S)H WCH2 WCH3

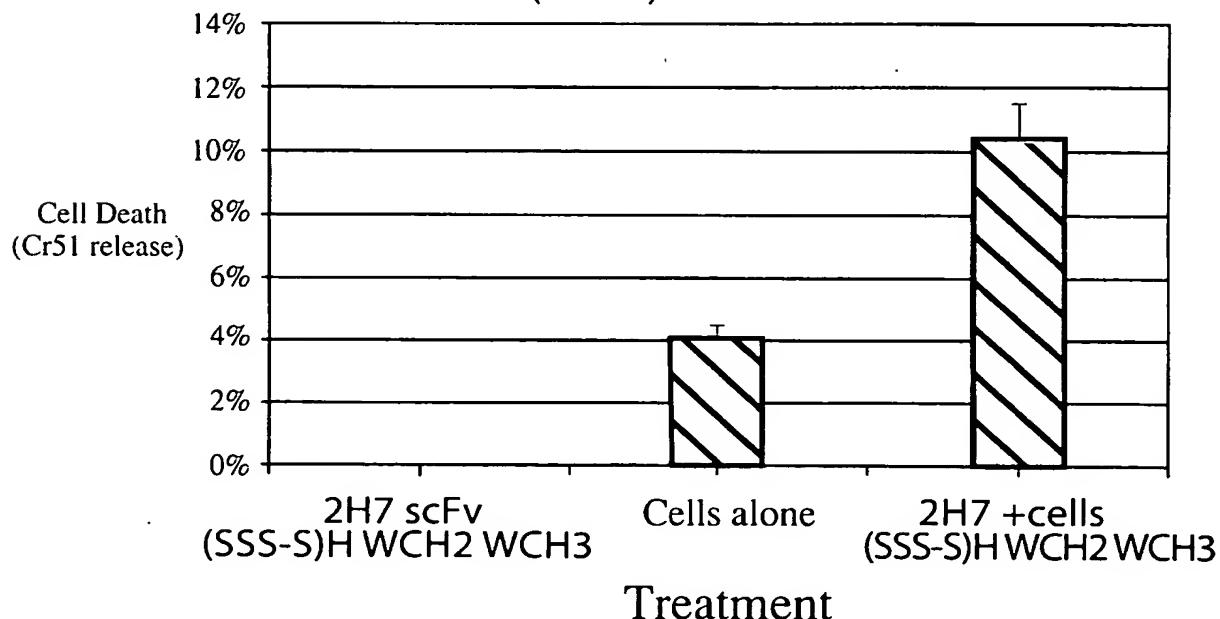


FIG. 5

Effects of Crosslinking of CD20 and CD40 Cell Surface Receptors on B Cell Proliferation:

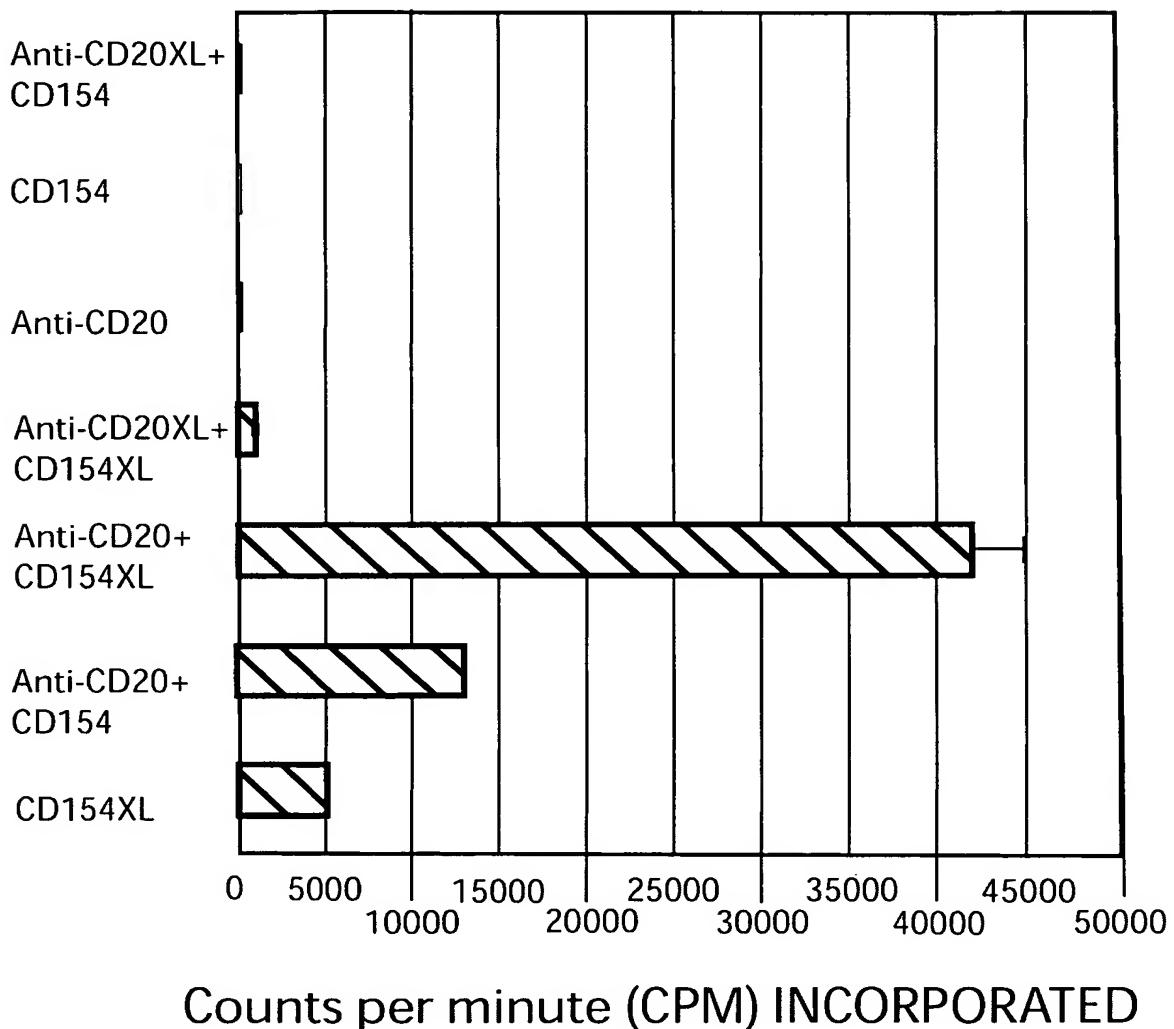
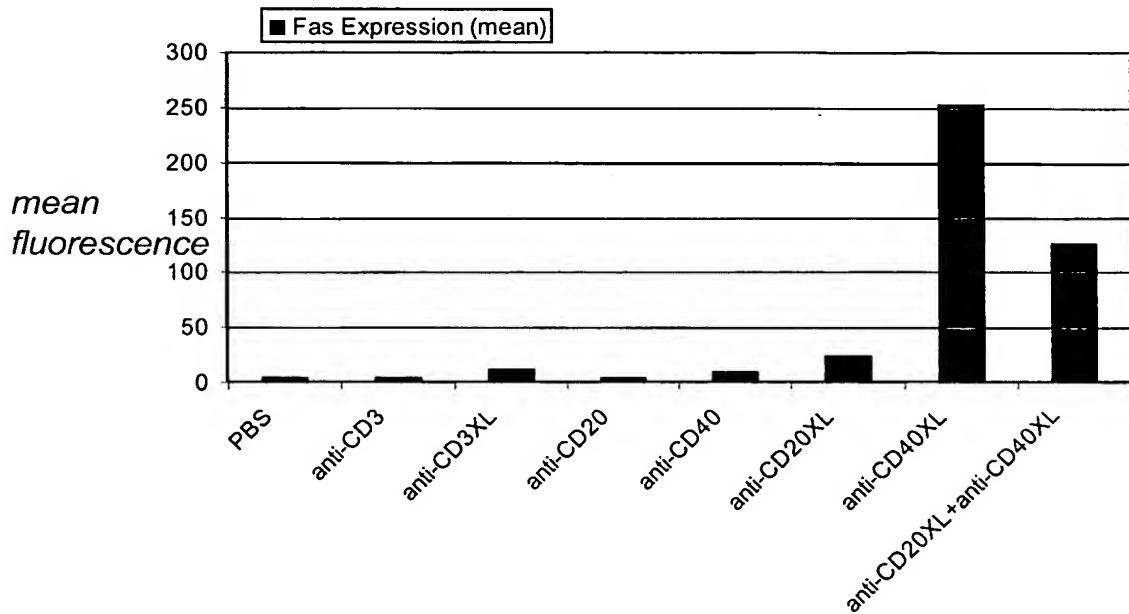


FIG. 6A

Effect of Simultaneous ligation of CD20 and CD40 on CD95 and apoptosis.

**FIG. 6B**

Effect of Simultaneous ligation of CD20 and CD40 on CD95 and apoptosis.

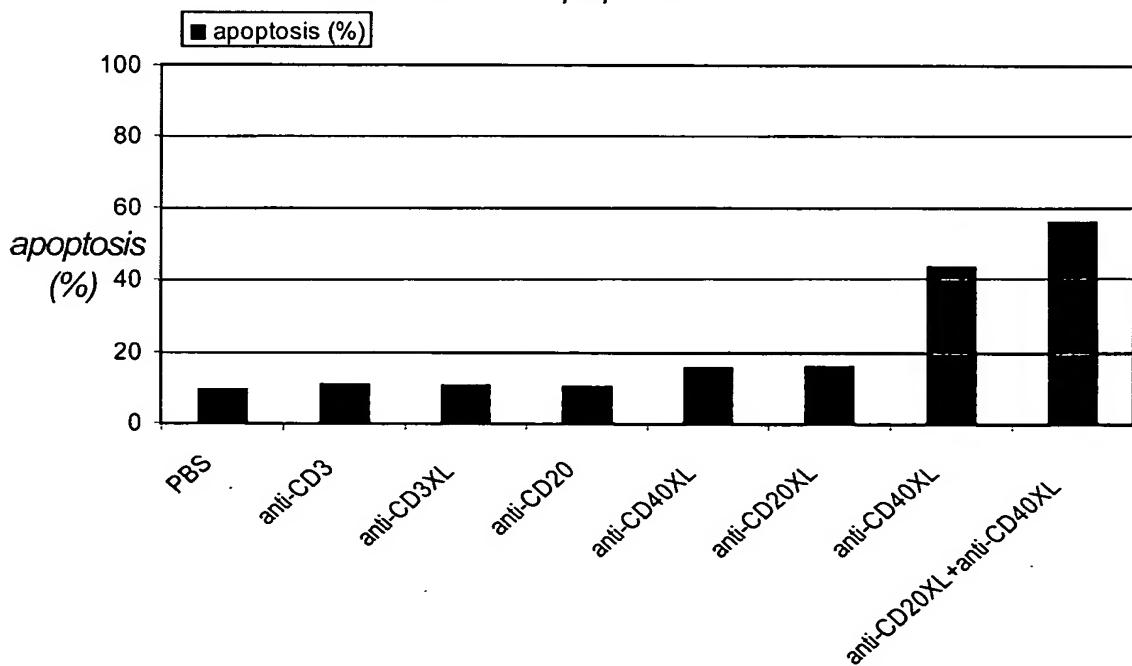


FIG. 7A

2H7-CD154 L2 cDNA and predicted amino acid sequence:

<pre>HindIII NcoI 2H7 V_L Leader Peptide → -----</pre>	<pre>1 M D F Q V Q I F S F L L I S A S AAGCTTGCCTG CC ATGGATTT TCAAGTGCAG ATTTTCAGCT TCCTGCTAAT CAGTGCTTCA</pre>
<pre>2H7 V_L → -----</pre>	<pre>61 V I I A R G Q I V L S Q S P A I L S A S GTCATAATTG CCAGAGGACA AATTGTTCTC TCCCAGTCTC CAGCAATCCT GTCTGCATCT</pre>
<pre>BamHI -----</pre>	<pre>121 P G E K V T M T C R A S S S V S Y M H W CCAGGGGAGA AGGTACAAAT GACTTCAGG GCCAGCTCAA GTGTAAGTTA CATGCACTGG</pre>
<pre>(Gly₄Ser)₃ Linker → -----</pre>	<pre>181 Y Q Q K P G S S P K P W I Y A P S N L A TACCAGCAGA AGCCAGGATC CTCCCCAAA CCCTGGATTT ATGCCCATC CAACCTGGCT</pre>
<pre>241 S G V P A R F S G S G S G T S Y S L T I TCTGGAGTCC CTGCTCGCTT CAGTGGCAGT GGGTCTGGGA CCTCTTACTC TCTCACAAATC</pre>	
<pre>301 S R V E A E D A A T Y Y C Q Q W S F N P AGCAGAGTGG AGGCTGAAGA TGCTGCCACT TATTACTGCC AGCAGTGGAG TTTAACCCA</pre>	
<pre>2H7 V_H → -----</pre>	<pre>361 P T F G A G T K L E L K G G G G S G G G CCCACGTTCG GTGCTGGAC CAAGCTGGAG CTGAAAGGTG GCGGTGGCTC GGGCGGTGGT</pre>
<pre>421 G S G G G G S S Q A Y L Q Q S G A E L V GGATCTGGAG GAGGTGGAG CTCTCAGGCT TATCTACAGC AGTCTGGGCG TGAGCTGGTG</pre>	
<pre>481 R P G A S V K M S C K A S G Y T F T S Y AGGCCTGGGG CCTCAGTGAAT GATGTCCTGC AAGGCTTCTG GCTACACATT TACCAAGTTAC</pre>	
<pre>541 N M H W V K Q T P R Q G L E W I G A I Y AATATGCAG GGGTAAAGCA GACACCTAGA CAGGGCCTGG AATGGATTGG AGCTATTAT</pre>	
<pre>601 P G N G D T S Y N Q K F K G K A T L T V CCAGGAAATG GTGATACTTC CTACAATCAG AAGTCAGG GCAAGGCCAC ACTGACTGTA</pre>	
<pre>661 D K S S S T A Y M Q L S S L T S E D S A GACAAATCCT CCAGCACAGC CTACATGCAG CTCAGCAGCC TGACATCTGA AGACTCTGCG</pre>	
<pre>721 V Y F C A R V V Y Y S N S Y W Y F D V W GTCTATTCT GTGCAAGAGT GGTGTACTAT AGTAACCTTT ACTGGTACTT CGATGTCTGG</pre>	

FIG. 7B

human CD154/amino acid 48→

site Bcl/Bam hybrid

G T G T T V T V S D P R R L D K I E D E
781 GGCACAGGGG CCACGGTCAC CGTCTCTGAT CCAAGAAGGT TGGACAAGAT AGAAGATGAA

R N L H E D F V F M K T I Q R C N T G E
841 AGGAATCTTC ATGAAGATTT TGTATTGATG AAAACGATAC AGAGATGCAA CACAGGAGAA

R S L S L L N C E E I K S Q F E G F V K
901 AGATCCTTAT CCTTACTGAA CTGTGAGGAG ATTAAAAGCC AGTTTGAGG CTTTGTGAAAG

BclI

D I M L N K E E T K K E N S F E M Q K G
961 GATATAATGT TAAACAAAGA GGAGACGAAG AAAGAAAACA GCTTTGAAAT GCAAAAAGGT

BcII

D Q N P Q I A A H V I S E A S S K T T S
1021 GATCAGAACATC CTCAAATTGC GGCACATGTC ATAAGTGAGG CCAGCAGTAA AACAAACATCT

V L Q W A E K G Y Y T M S N N L V T L E
1081 GTGTTACAGT GGGCTGAAAA AGGATACTAC ACCATGAGCA ACAACTTGTT AACCTGGAA

N G K Q L T V K R Q G L Y Y I Y A Q V T
1141 AATGGGAAAC AGCTGACCGT TAAAAGACAA GGACTCTATT ATATCTATGC CCAAGTCACC

HindIII

F C S N R E A S S Q A P F I A S L C L K
1201 TTCTGTTCCA ATCGGGAAAGC TTCGAGTCAA GCTCCATTAA TAGCCAGCCT CTGCCTAAAG

S P G R F E R I L L R A A N T H S S A K
1261 TCCCCCGGTA GATTGAGAG AATCTTACTC AGAGCTGCAA ATACCCACAG TTCCGCCAAA

P C G Q Q S I H L G G V F E L Q P G A S
1321 CCTTGCGGGC AACAAATCCAT TCACTTGGGA GGAGTATTTG AATTGCAACC AGGTGCTTCG

NcoI

V F V N V T D P S Q V S H G T G F T S F
1381 GTGTTGTCA ATGTGACTGA TCCAAGCCAA GTGAGCCATG GCACTGGCTT CACGTCCTTT

XbaI

G L L K L E * * S R
1441 GGCTTACTCA AACTCGAGTG ATAATCTAGA

FIG. 7C

2H7scFv-CD154 S4 cDNA and predicted amino acid sequence:

HindIII NcoI
 ----- 2H7 V_L Leader Peptide →
 M D F Q V Q I F S F L L I S A S
 1 AAGCTTGCCG CC ATGGATTT TCAAAGTGCAG ATTTTCAGCT TCCTGCTAAT CAGTGCTTCA

2H7 V_L →
 V I I A R G Q I V L S Q S P A I L S A S
 61 GTCATAATTG CCAGAGGACA AATTGTTCTC TCCCAGTCTC CAGCAATCCT GTCTGCATCT

P G E K V T M T C R A S S S V S Y M H W
 121 CCAGGGGAGA AGGTCAACAAT GACTTGCAGG GCCAGCTCAA GTGTAAGTTA CATGCACTGG

BamHI

 Y Q Q K P G S S P K P W I Y A P S N L A
 181 TACCAAGCAGA AGCCAGGATC CTCCCCCAA CCCTGGATTG ATGCCCATC CAACCTGGCT

S G V P A R F S G S G S G T S Y S L T I
 241 TCTGGAGTCC CTGCTCGCTT CAGTGGCAGT GGGTCTGGGA CCTCTTACTC TCTCACAACTC

S R V E A E D A A T Y Y C Q Q W S F N P
 301 AGCAGAGTGG AGGCTGAAGA TGCTGCCACT TATTACTGCC AGCAGTGGAG TTTTAACCCA

(Gly₄Ser)₃ Linker →
 P T F G A G T K L E L K G G G G S G G G
 361 CCCACGTTCG GTGCTGGAC CAAGCTGGAG CTGAAAGGTG GCGGTGGCTC GGGCGGTGGT

2H7 V_H →
 G S G G G G S S Q A Y L Q Q S G A E L V
 421 GGATCTGGAG GAGGTGGGAG CTCTCAGGCT TATCTACAGC AGTCTGGGC TGAGCTGGTG

R P G A S V K M S C K A S G Y T F T S Y
 481 AGGCCTGGGG CCTCAAGTCAA GATGTCTGC AAGGCTTCTG GCTACACATT TACCAGTTAC

N M H W V K Q T P R Q G L E W I G A I Y
 541 AATATGCACT GGGTAAAGCA GACACCTAGA CAGGGCCTGG AATGGATTGG AGCTATTAT

P G N G D T S Y N Q K F K G K A T L T V
 601 CCAGGAAATG GTGATACTTC CTACAATCAG AAGTTCAAGG GCAAGGCCAC ACTGACTGTA

D K S S S T A Y M Q L S S L T S E D S A
 661 GACAAATCCT CCAGCACAGC CTACATGCAG CTCAGCAGCC TGACATCTGA AGACTCTGCG

V Y F C A R V V Y Y S N S Y W Y F D V W
 721 GTCTATTTCT GTGCAAGAGT GGTGTACTAT AGTAACCTTT ACTGGTACTT CGATGTCTGG

FIG.7D

human CD154/amino acid 108 →

BclI Bcl/Bam hybrid site

781 G T G T T V T V S D P E N S F E M Q K G
GGCACAGGGGA CCACGGTCAC CGTCTCTGAT CCAGAAAACA GCTTGAAAT GCAAAAGGT

BclI

841 D Q N P Q I A A H V I S E A S S K T T S
GATCAGAACATC CTCAAATTGC GGCACATGTC ATAAGTGAGG CCAGCAGTAA AACAAACATCT

901 V L Q W A E K G Y Y T M S N N L V T L E
GTGTTACAGT GGGCTGAAAA AGGATACTAC ACCATGAGCA ACAACTTGGT AACCCCTGGAA

961 N G K Q L T V K R Q G L Y Y I Y A Q V T
AATGGGAAAC AGCTGACCGT TAAAAGACAA GGACTCTATT ATATCTATGC CCAAGTCACC

HindIII

1021 F C S N R E A S S Q A P F I A S L C L K
TTCTGTTCCA ATCGGGAAAGC TTCGAGTCAA GCTCCATTAA TAGCCAGCCT CTGCCCTAAAG

1081 S P G R F E R I L L R A A N T H S S A K
TCCCCCGGTA GATTGAGAG AATCTTACTC AGAGCTGCAA ATACCCACAG TTCCGCCAAA

1141 P C G Q Q S I H L G G V F E L Q P G A S
CCTTGCAGGC AACAAATCCAT TCACTTGGGA GGAGTATTTG AATTGCAACC AGGTGCTTCG

NcoI

1201 V F V N V T D P S Q V S H G T G F T S F
GTGTTGTCA ATGTGACTGA TCCAAGCCAA GTGAGCCATG GCACTGGCTT CACGTCCCTT

XbaI

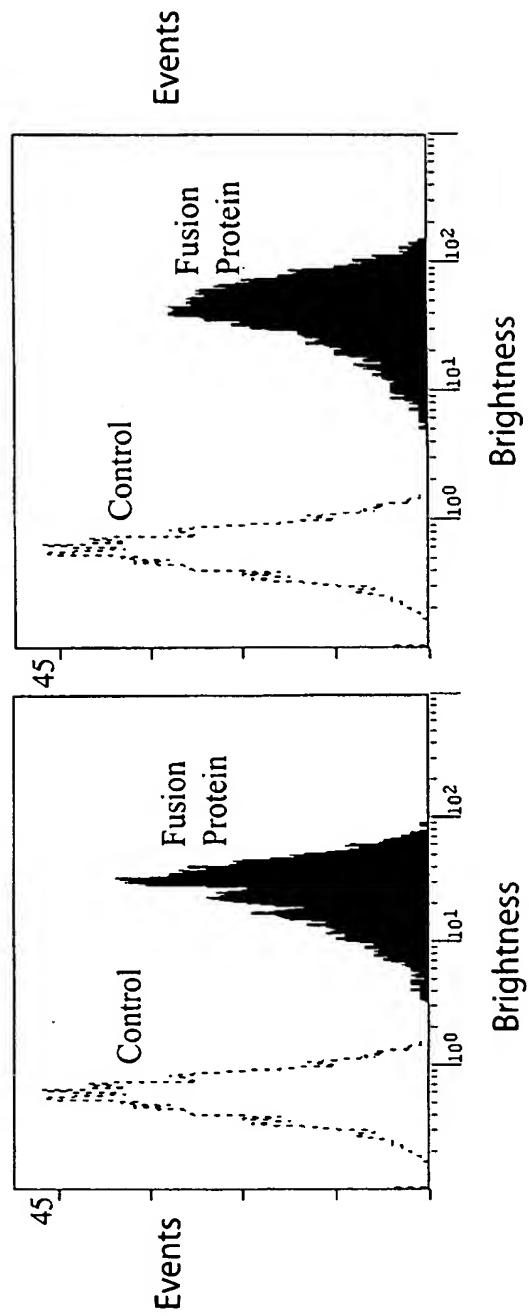
1261 G L L K L E * * S R
GGCTTACTCA AACTCGAGTG ATAATCTAGA

FIG. 8

Simultaneous Binding of 2H7scFv-CD154 Fusion Proteins to CD20 and CD40

2H7scFv-CD154
Construct S4

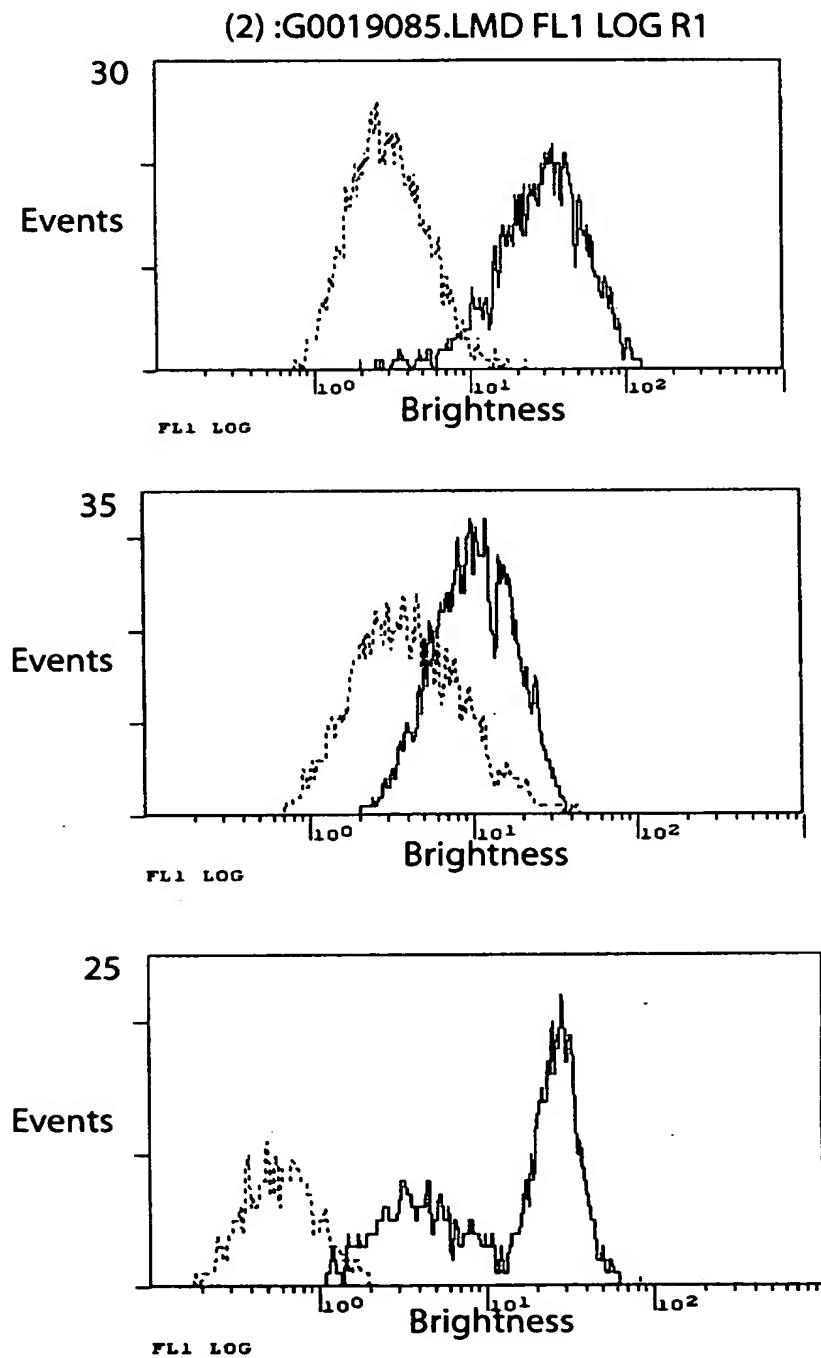
2H7scFv-CD154
Construct L2



CD20 CHO cell targets + (control or fusion protein)
+ Biotin-CD40Ig + PE-SA

FIG.9

Induction of Apoptosis Measured by Binding of Annexin V after incubation with 2H7scFv-CD154



.....control supernatant 2H7scFv-CD154 supernatant

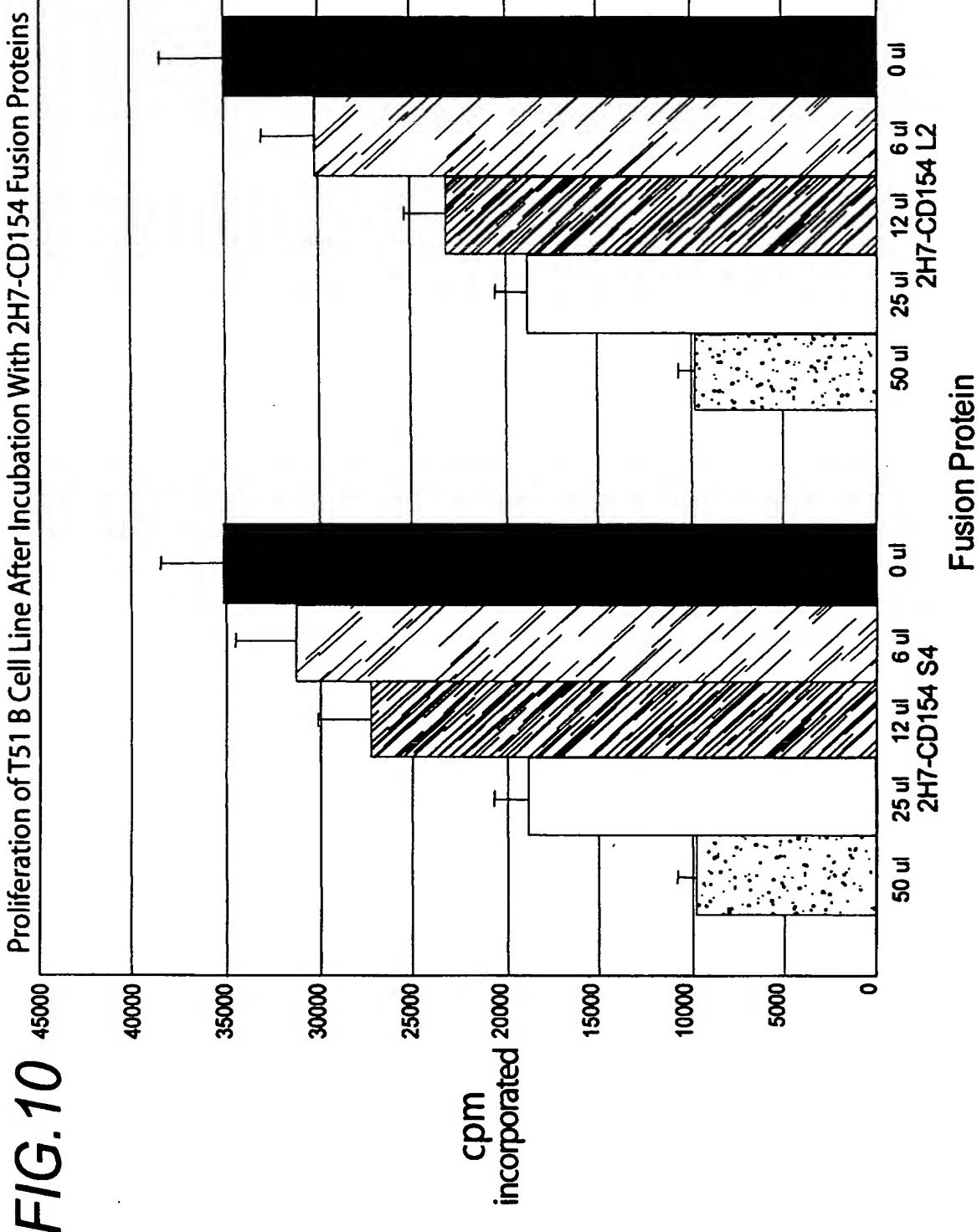


FIG. 11

2H7 scFv (SSS-S)H WCH2 WCH3
OR 2H7 scFv (SSS-S)H P238SCH2 WCH3

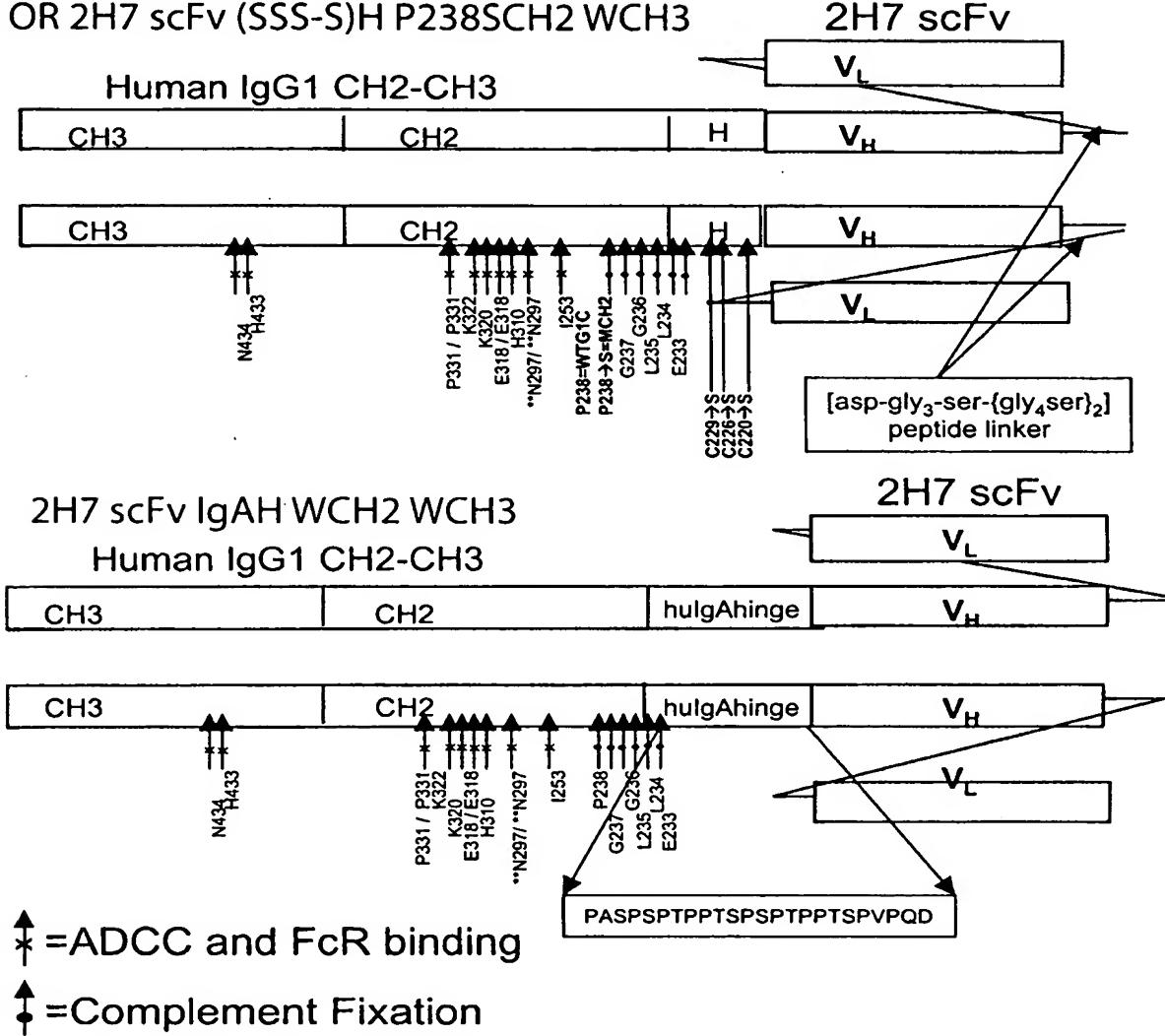


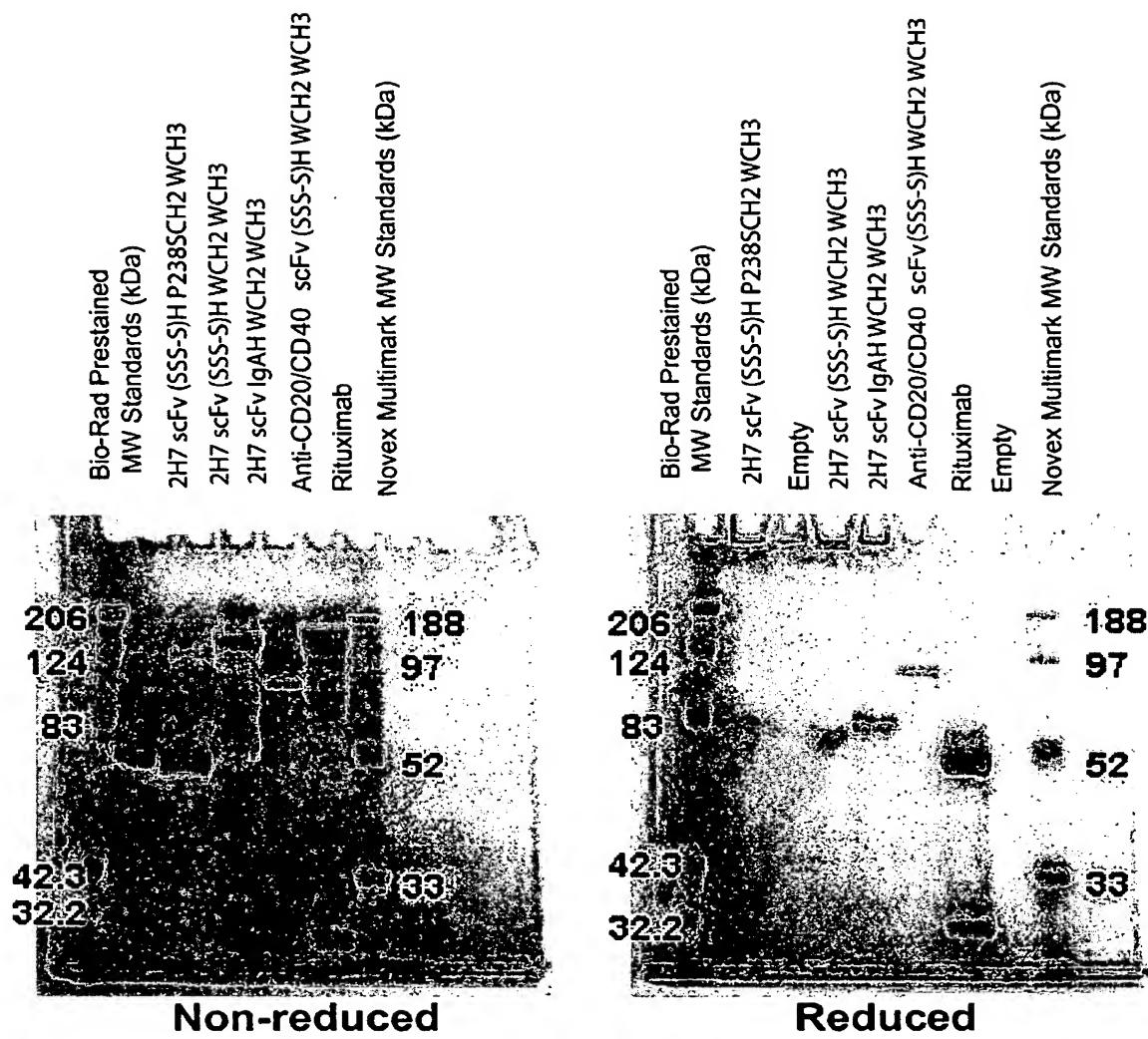
FIG. 12

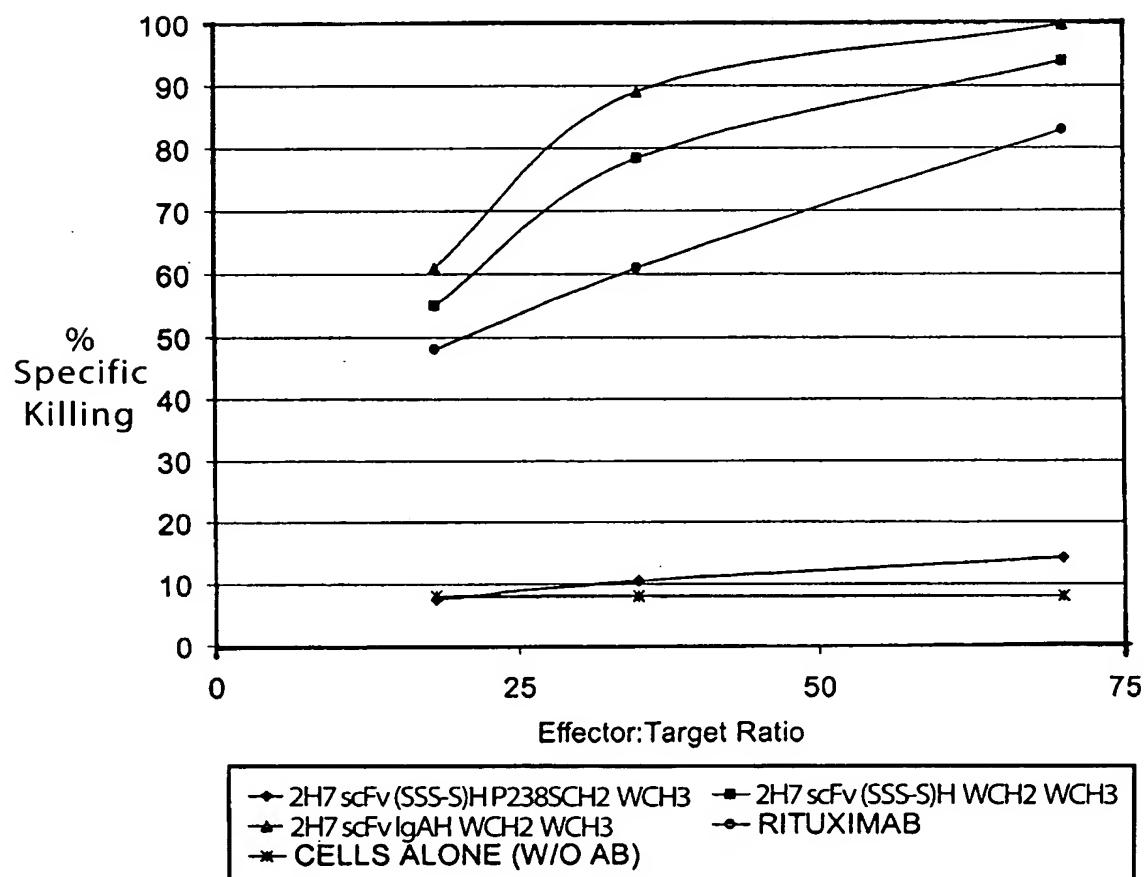
FIG. 13**ADCC Activity of 2H7 scFv Constructs**

FIG. 14

18/75

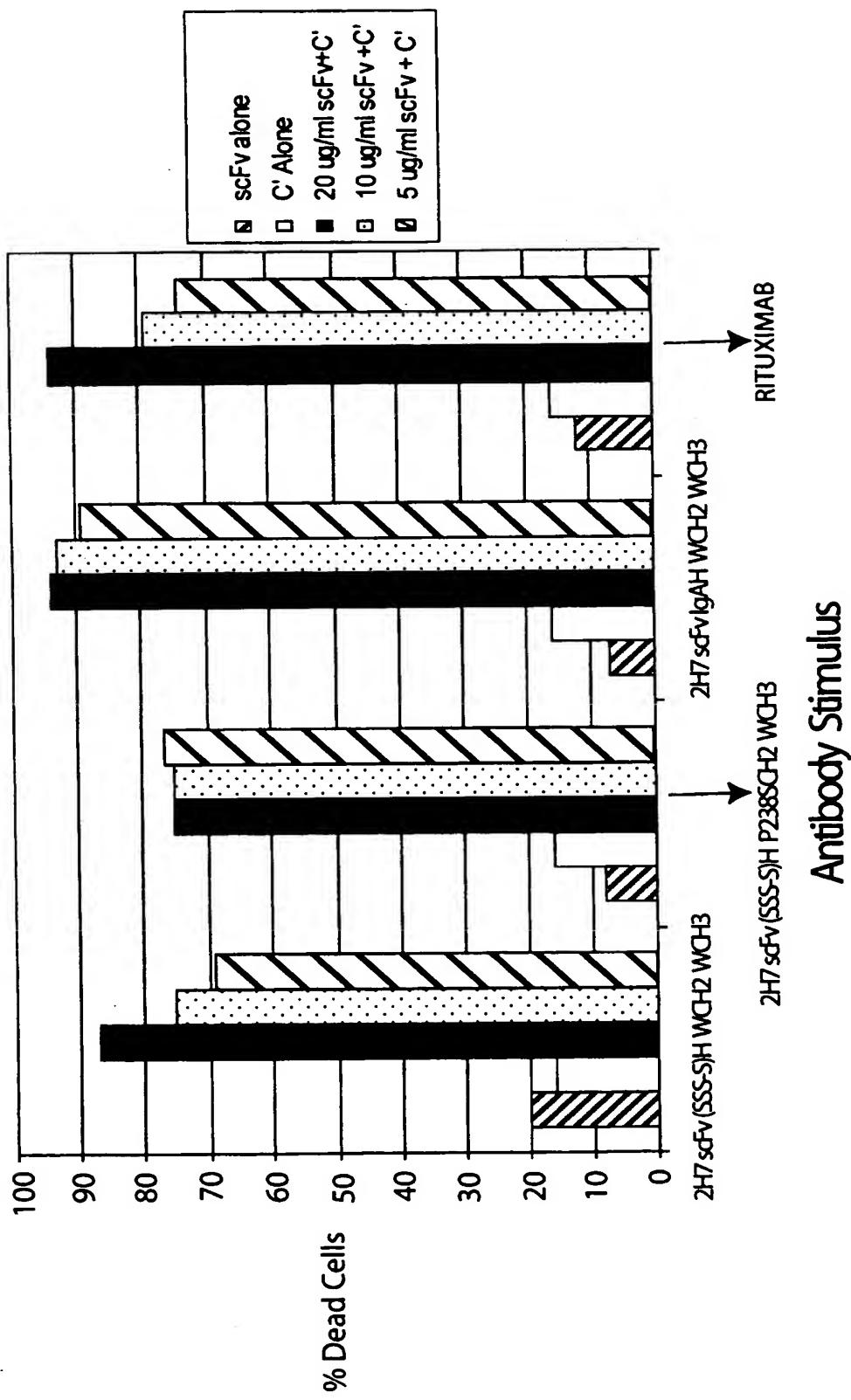
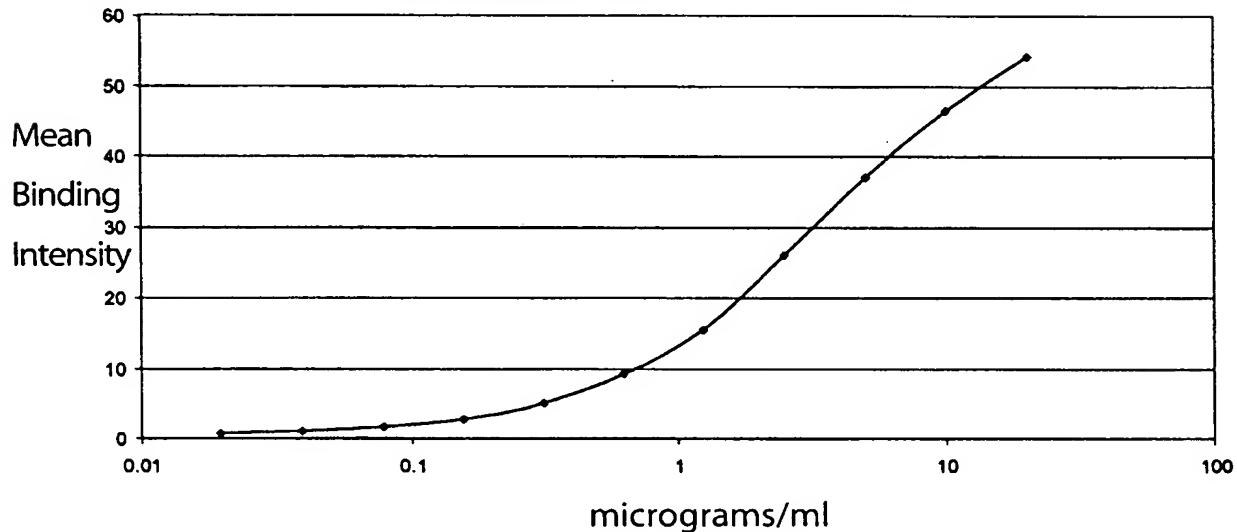


FIG. 15**2H7 scFv (SSS-S)H WCH2 WCH3 In Vivo Half Life****2H7 scFv (SSS-S)H WCH2 WCH3 Standard Curve****Macaque A99314**

	<u>Day</u>	Binding intensity At 1:50	estimated concentration ($\mu\text{g}/\text{ml}$)
Injection #1 →	-7	0.213	<0.1
	0	0.227	<0.1
	1	7.79	25.1
	3	5.51	15.6
Injection #2 →	7	3.37	9.4
	8	11.33	41.7
	10	5.45	15.4
	14	0.27	<0.1

Macaque F98081

	<u>Day</u>	Binding intensity At 1:50	estimated concentration ($\mu\text{g}/\text{ml}$)
Injection #1 →	-7	0.208	<0.1
	0	0.219	<0.1
	1	6.73	21.9
	3	6.14	19.3
Injection #2 →	7	3.04	8.7
	8	9.83	33.8
	10	4.77	14.4
	14	0.231	<0.1

FIG. 16

B Cell Depletion in macaques mediated by
2H7 scFv (SSS-S)H WCH2 WCH3

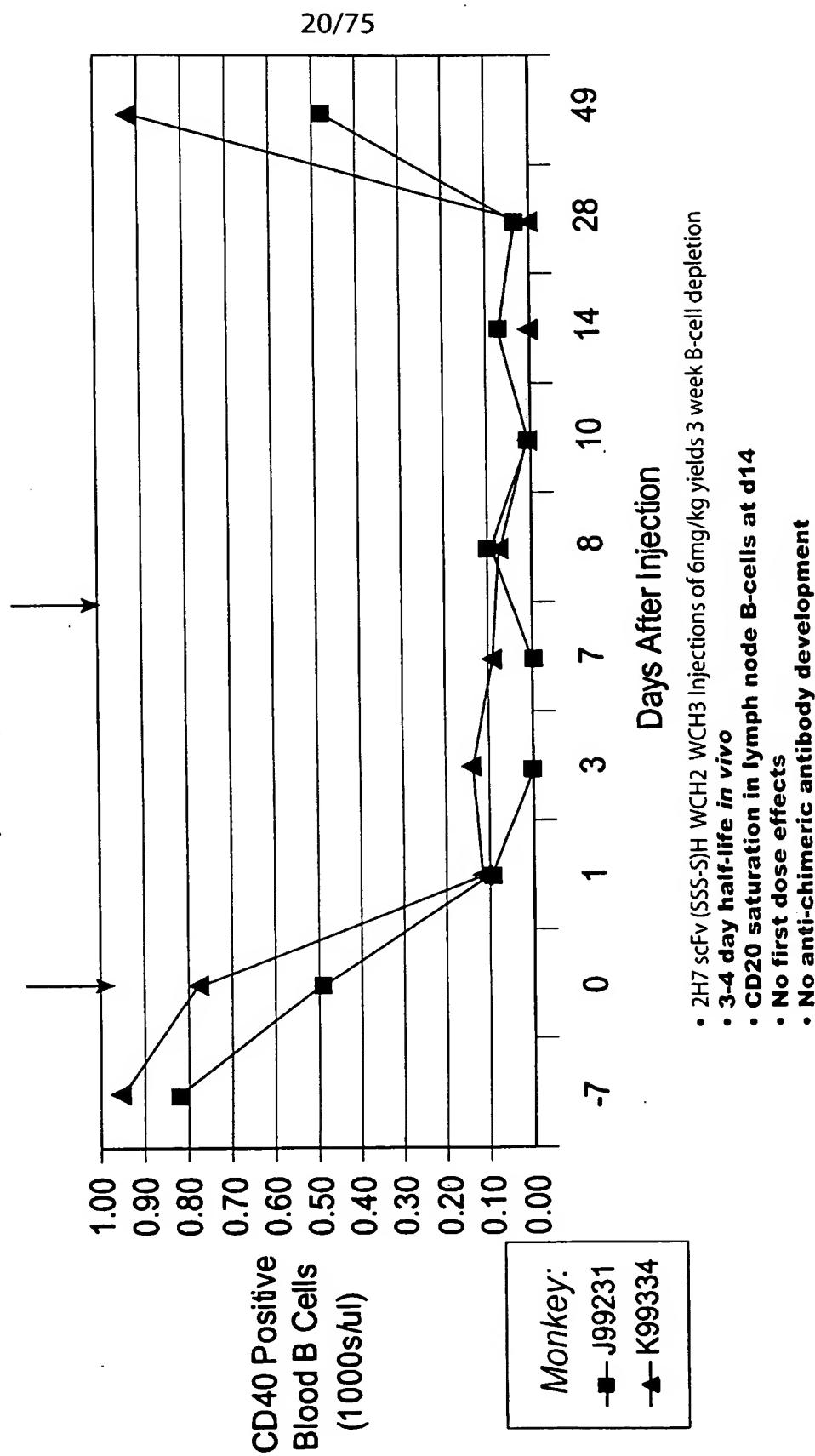
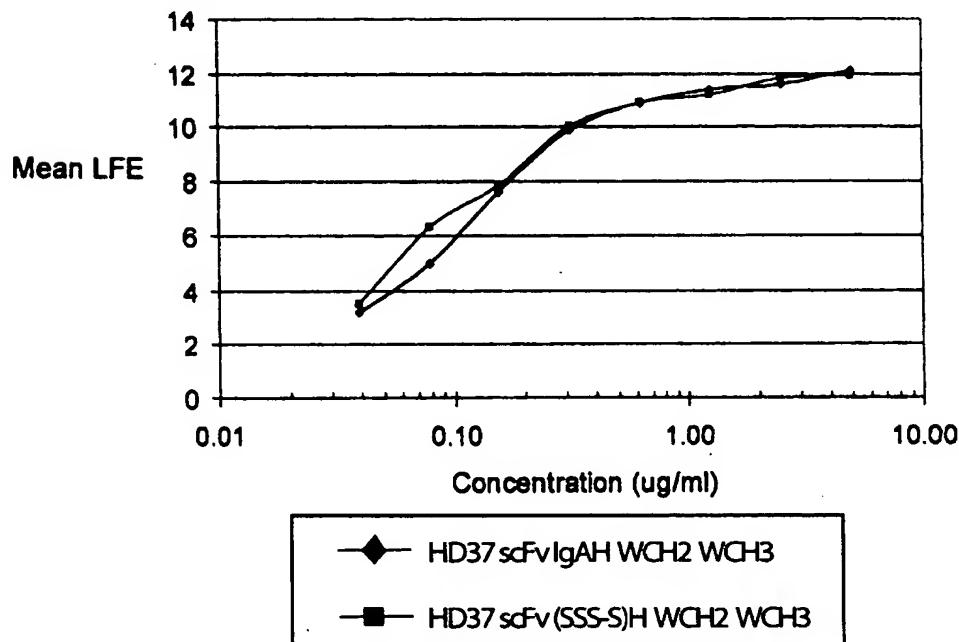


FIG. 17**Production Levels of HD37 scFv Constructs by CHO Cell Lines****Standard Curve of HD37 scFvIg Constructs
Binding to B Cells**

<u>Clone/Isolate</u>	<u>Mean LFE at 1:100</u>	<u>Estimated Concentration</u>
HD37 scFvIgAH WCH2 WCH3	11.2	> 60 ug/ml
1B2	10.4	>50 ug/ml
6C5	10.5	>50 ug/ml
4B1	8.6	>40 ug/ml
HD37 scFv(SSS-S)H WCH2 WCH3	10.9	> 50 ug/ml
2G8	10.6	> 50 ug/ml
3F3	8.3	>40 ug/ml
3D9	11.1	> 60 ug/ml

FIG. 18 Production of L6 scFvlg Constructs by CHO Cells

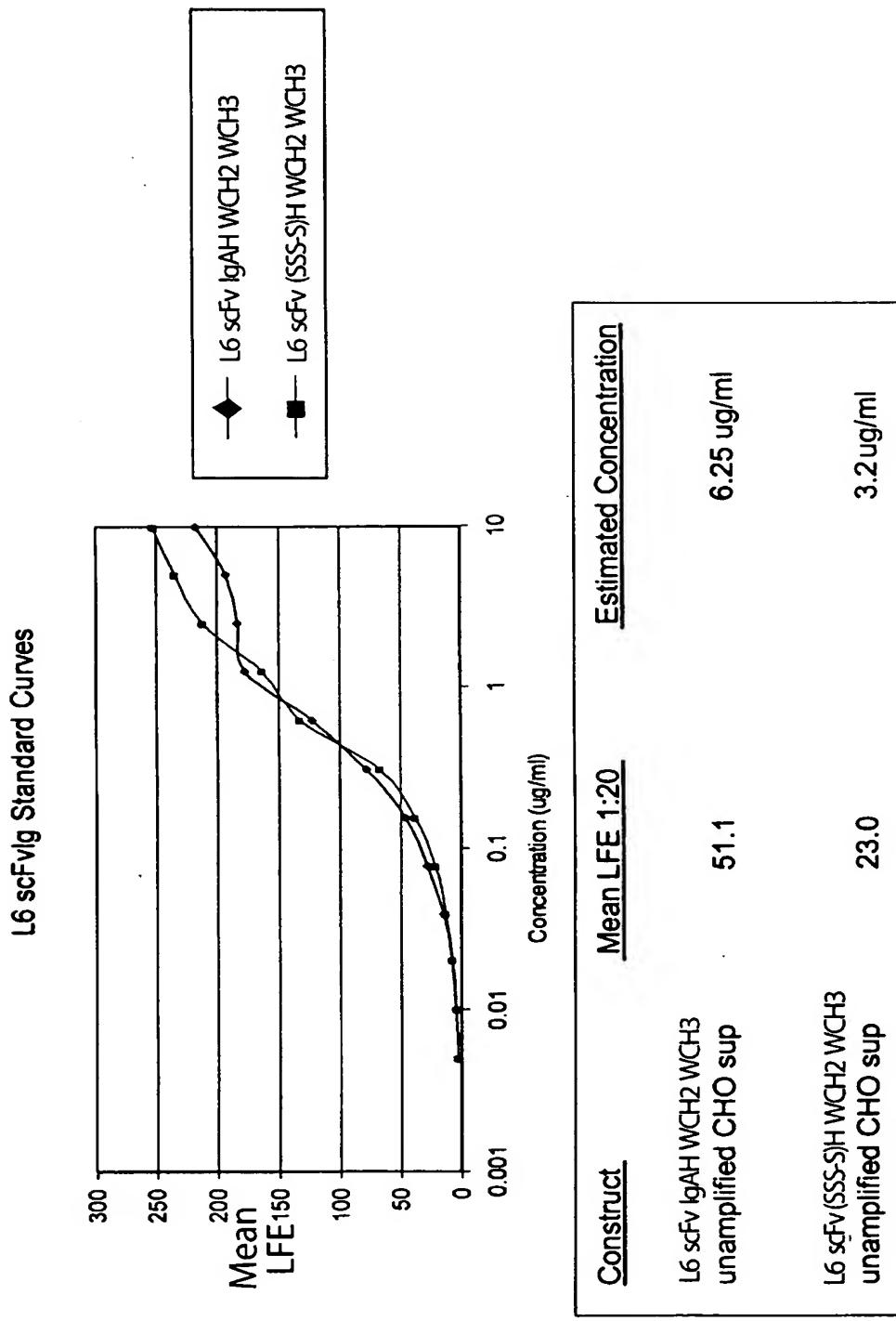


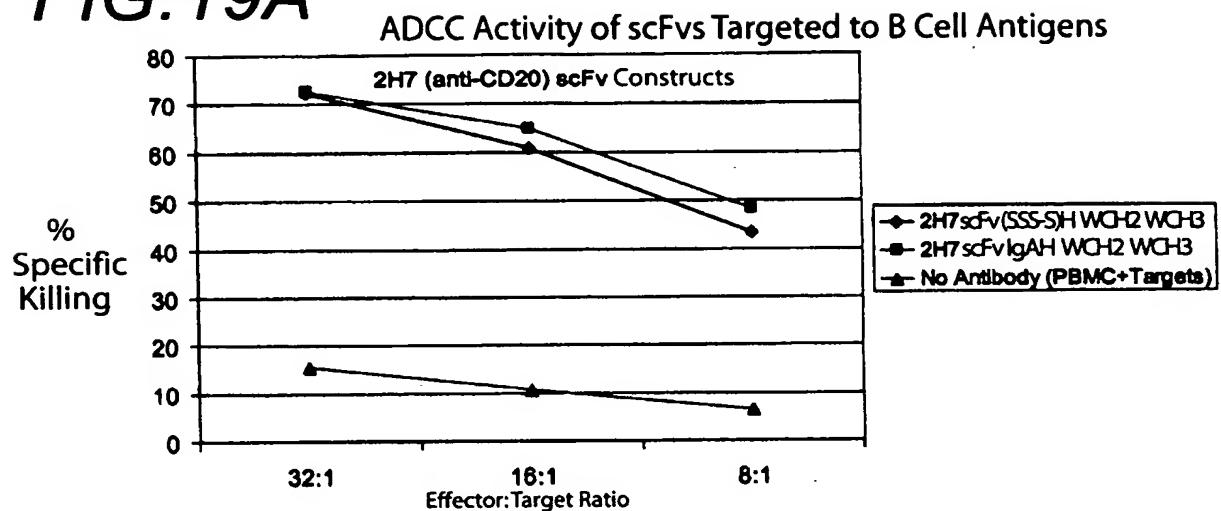
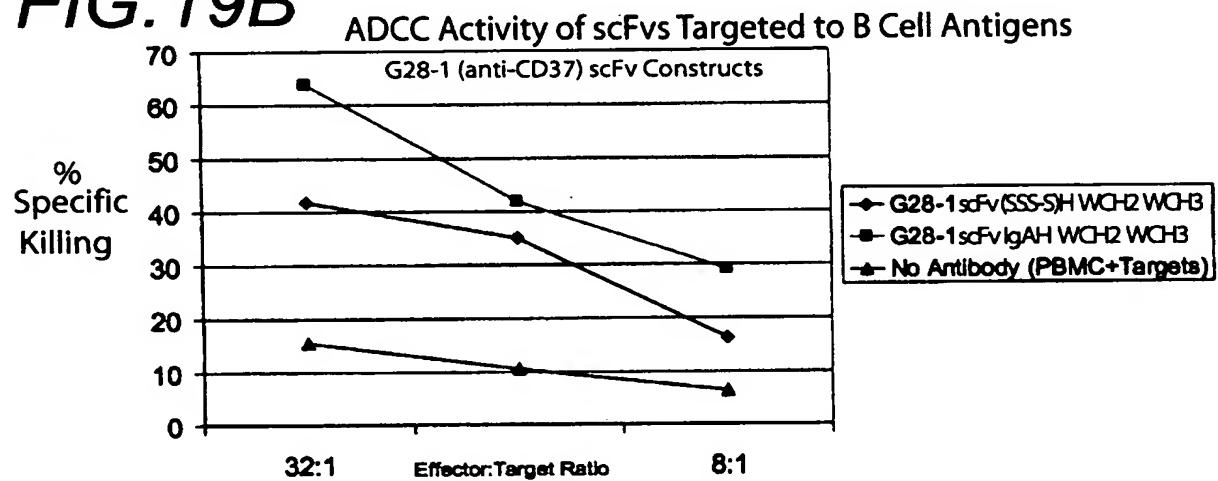
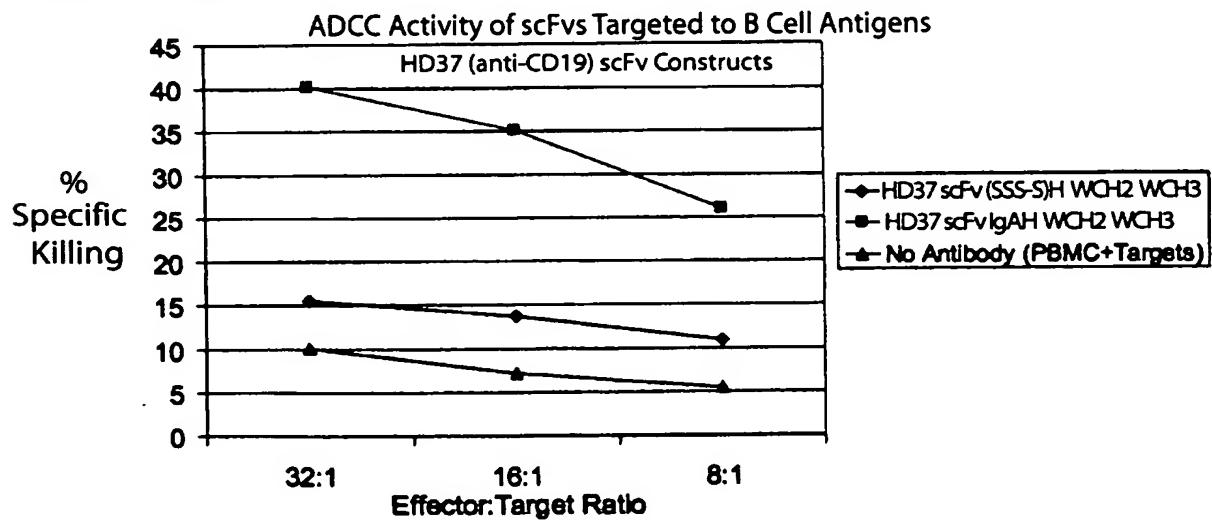
FIG. 19A**FIG. 19B****FIG. 19C**

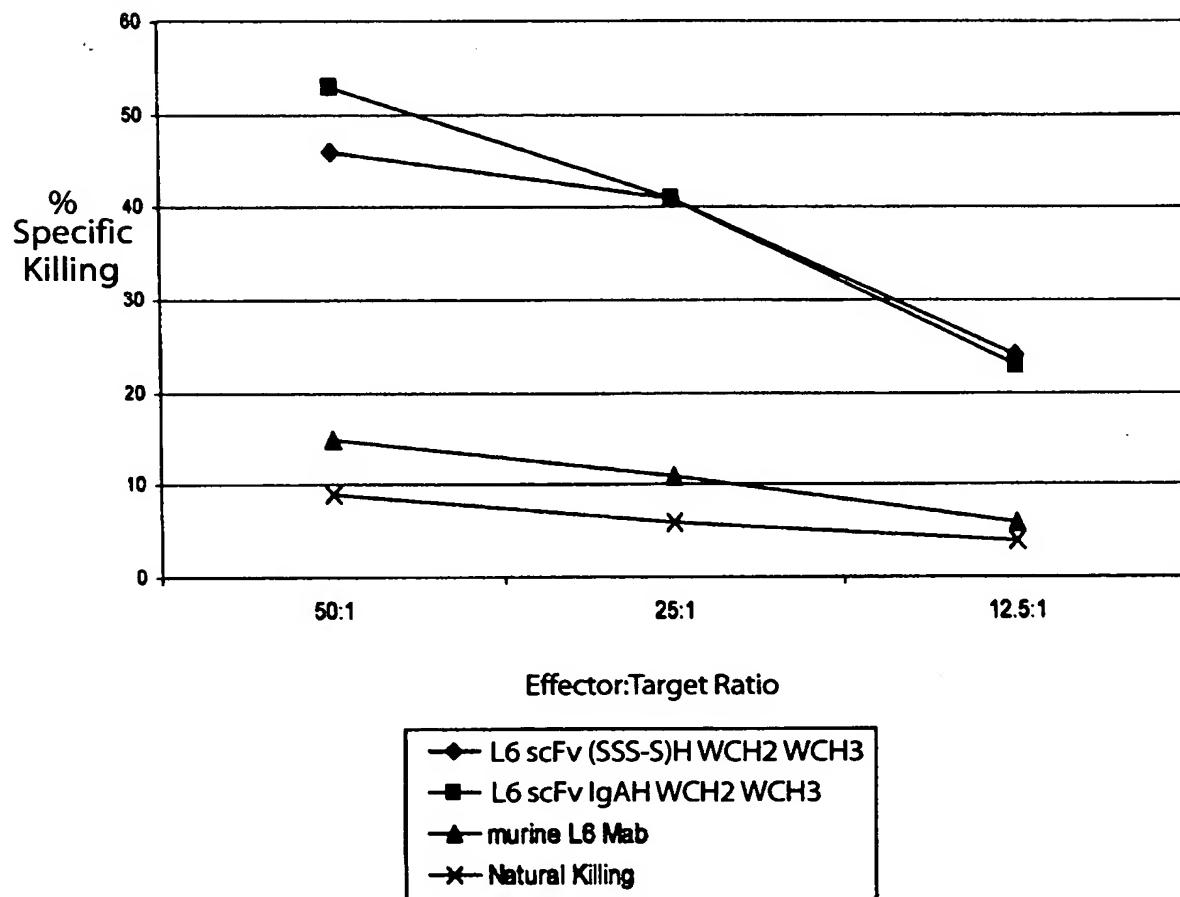
FIG. 20**ADCC Activity of L6 scFvIg Constructs****ADCC Activity of L6scFvIg Constructs with 2981 Targets**

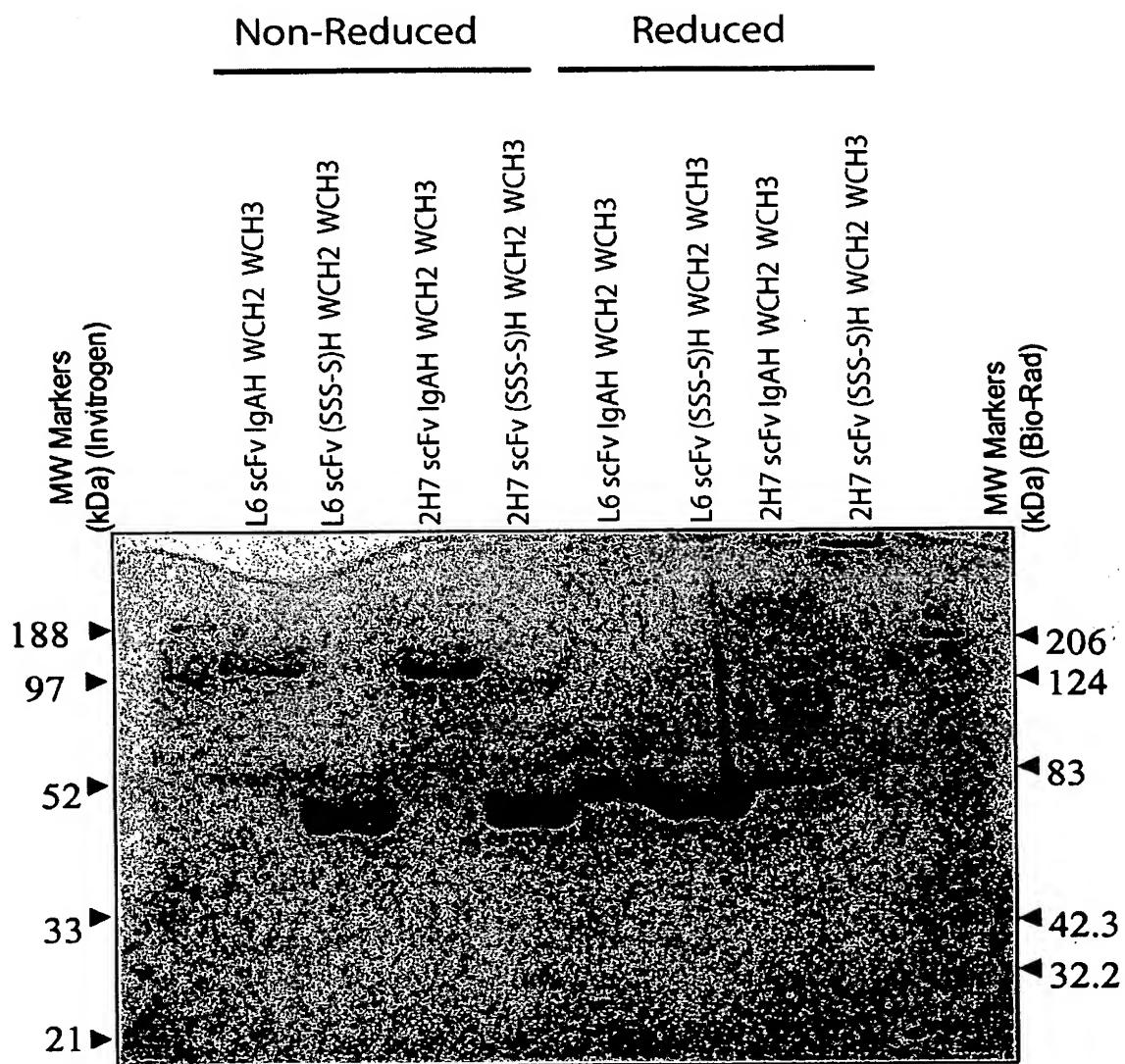
FIG. 21SDS-PAGE Analysis of L6 and 2H7
scFvIg Constructs

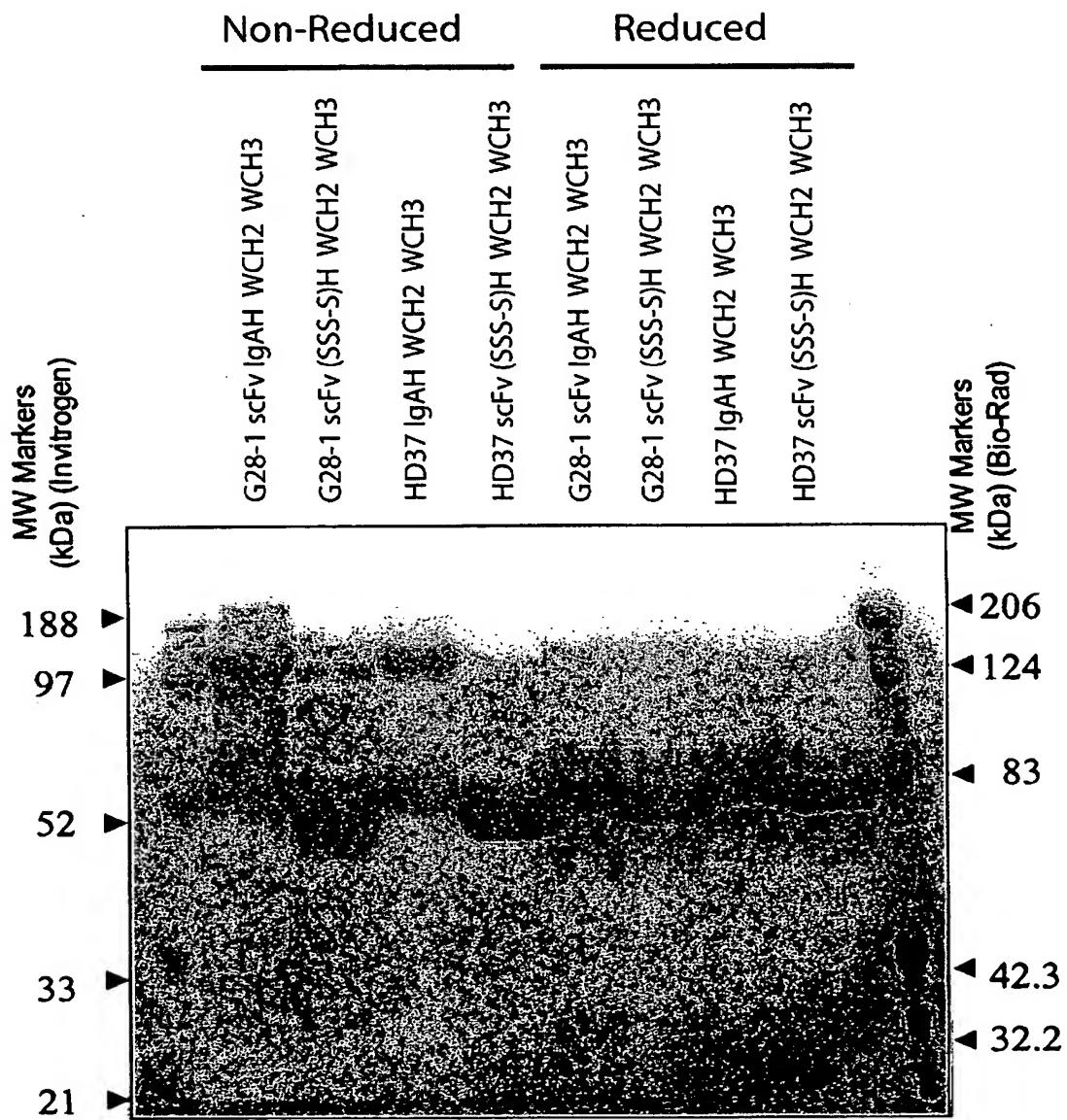
FIG.22**SDS-PAGE Analysis of G28-1 and HD37 scFvIg Constructs**

FIG. 23

Sequence alignment of human and Llama Fc regins.

HINGE CH2 →

Human IgG1:	DQEPKSCDKT-----	HTCPPC	PAPELLGGPSVFLFPKKPKDITLMISRTPEVTCAVVDVSHEDPEVKFNWYVDG
Llama IgG2:	DQEPKTPKPKPQPOQPPQPNPTTESKCPKC	PAPELLGGPSVFIFFPKPKDVLISIGRPEVTCAVVDVGQEDPEVSFNWYIDG	
Llama IgG1:	--EPHGG-----	CTCPQC	PAPELLGGPSVFFPKPKDVLISIGRPEVTCAVVDVGKEDPEVNFMNWFIDG
Llama IgG3:	--AHHSEDP-----	SKCPKO	PGPELLGGPPTVFIFFPKAKDVLISITRKPEVTCLWWTWVKKTLRSSSSWSVDD

VEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREGQQVYTLPSSRDELTKNQVSILT
TAEVRANTRPKKEQFNSTYRVVSVLPIQHQDWLTGKEFKCKVNNKALPAPIEKTISKAKGQTREPQVYTLAPFHREELAKDTVSVT
VEVRTANTKPKKEQFNSTYRVVSVLPIQHQDWLTGKEFKCKVNNKALPAPIERTISKAKGQTREPQVYTLAPFHREELAKDTVSVT
TEVHTAETKPKKEQFNSTYRVVSVLPIQHQDWLTGKEFKCKVNNKALPAPIERTISKAKGQTREPQVYTLAPFHREELAKDTVSVT

CLVKGFYPSDIAVEWESNGOPEN--NYKTTTPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSYMHEALHNHYTOKSLSLSPGK
CLVKGFYPPDINNEWQRNGQPESEGTYANTPPQLDNDGTYFLXSKXSVGKNTWQQGETFTCVVMHEALHNHYTOKSITQSSGK
CLVKGFYPADINNEWQRNGQPESEGTYANTPPQLDNDGTYFLYSRLSVGKNTWQRGETLTGVVMHEALHNHYTOKSITQSSGK
CLVKGFFPADINNEWQRNGQPESEGTYANTPPQLDNDGTYFLYSKLSVGKNTWQQGEVFTCVVMHEALHNHSTQKSITQSSGK

FIG. 24

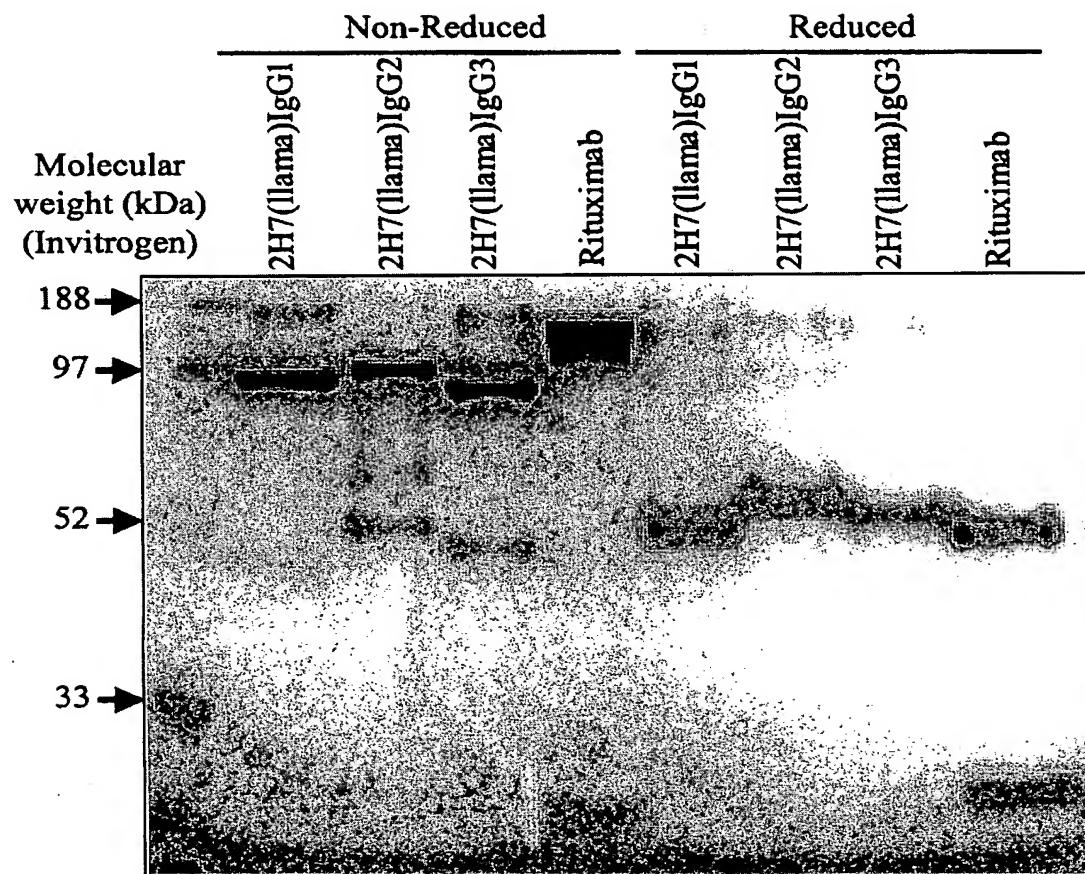
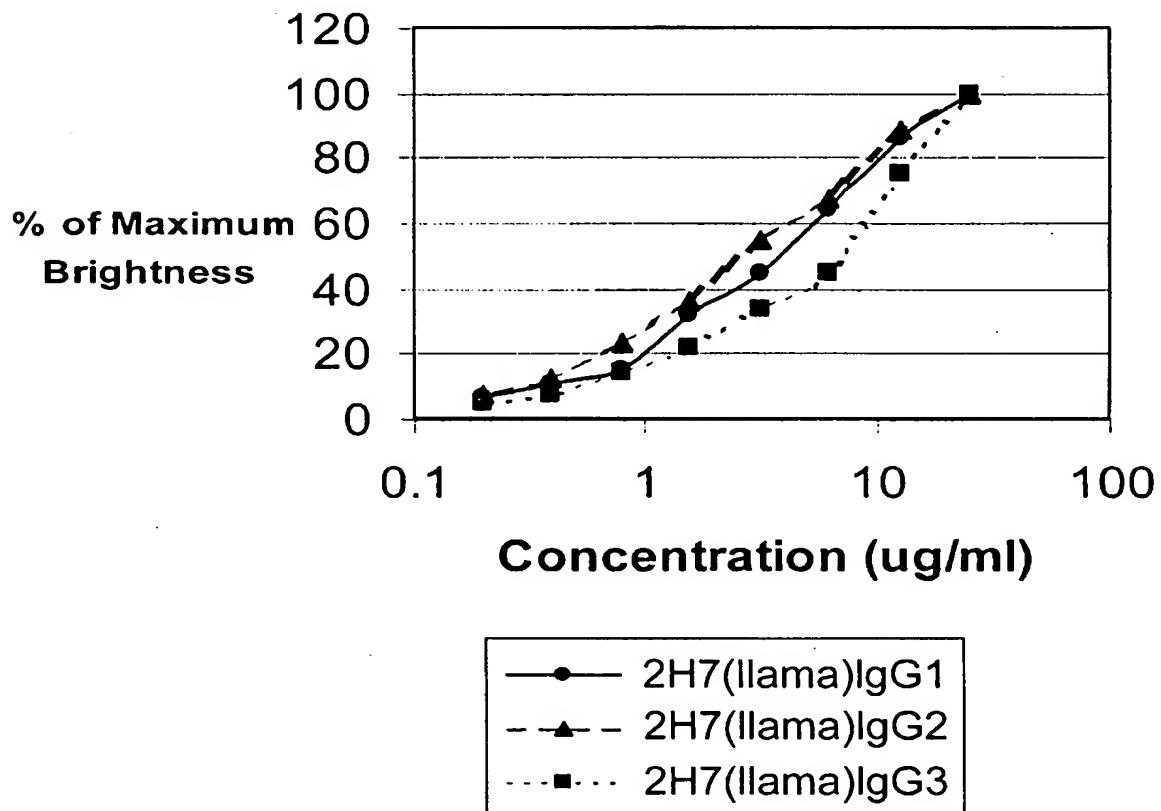


FIG.25**Llama Tails Binding Assay with CD20 CHO Cells**

30/75

FIG.26

2H7 scFv Llama Constructs Complement Assay with BJAB Cells

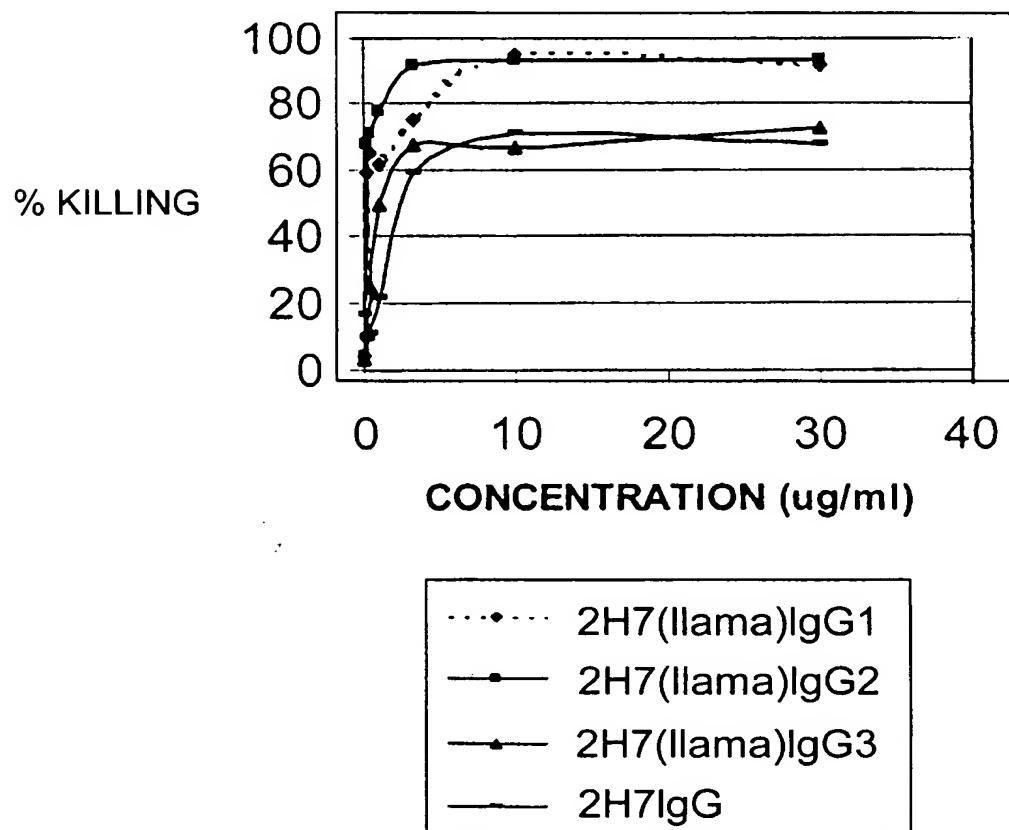


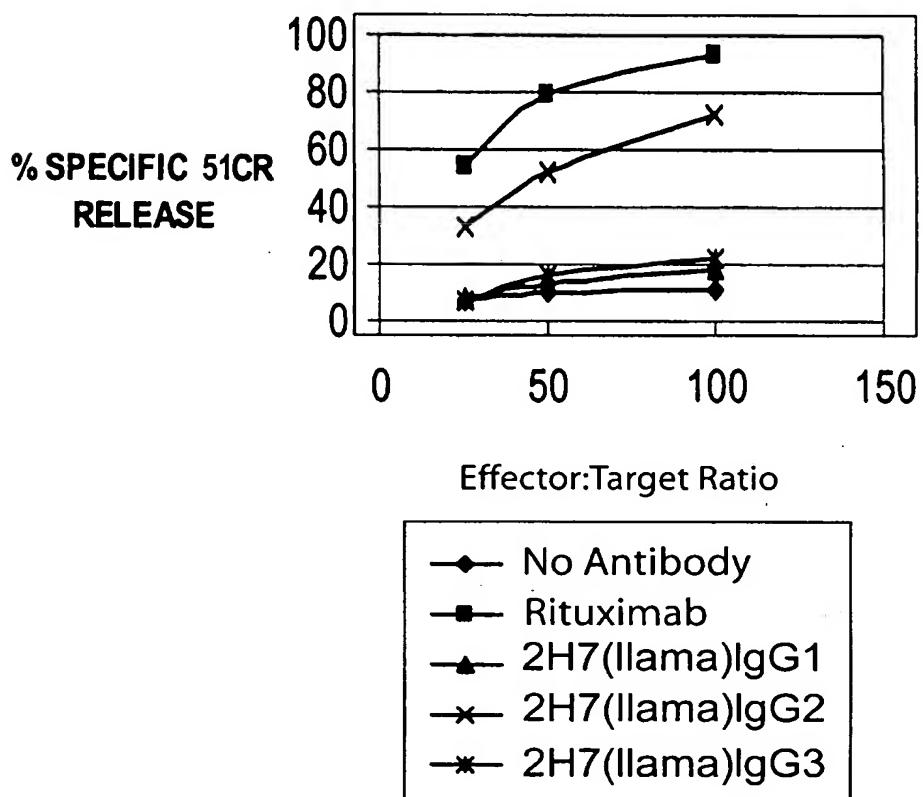
FIG.27**ADCC ASSAY WITH BJAB TARGETS
AND HUMAN PBMC EFFECTORS**

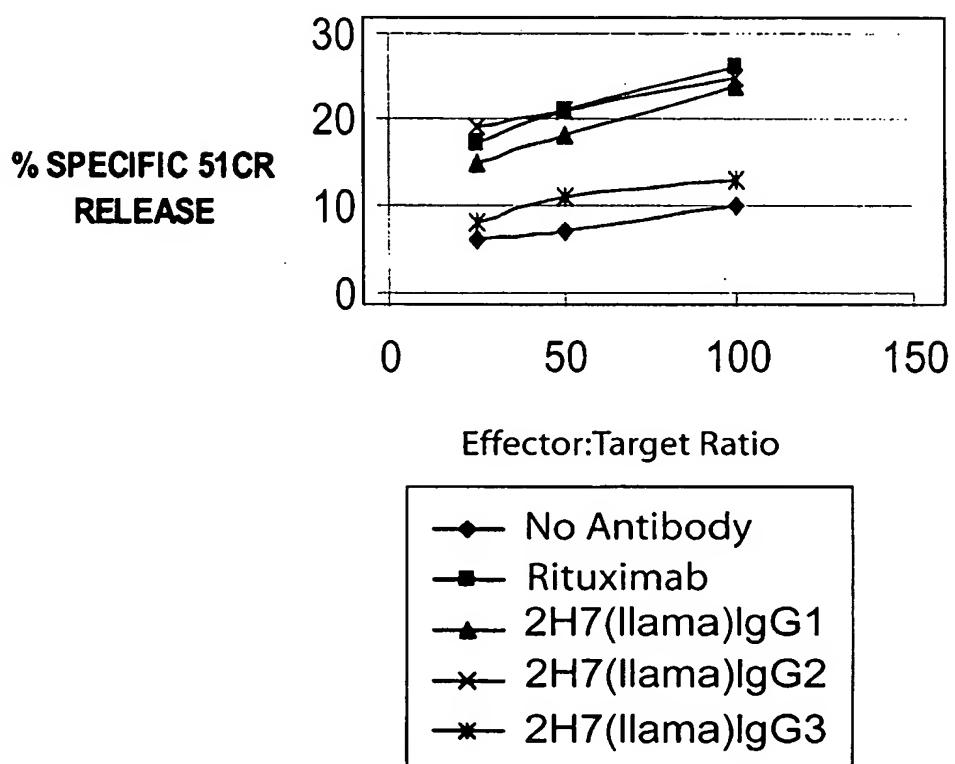
FIG.28ADCC ASSAY WITH BJAB TARGETS
AND LLAMA PBMC EFFECTORS

FIG. 29

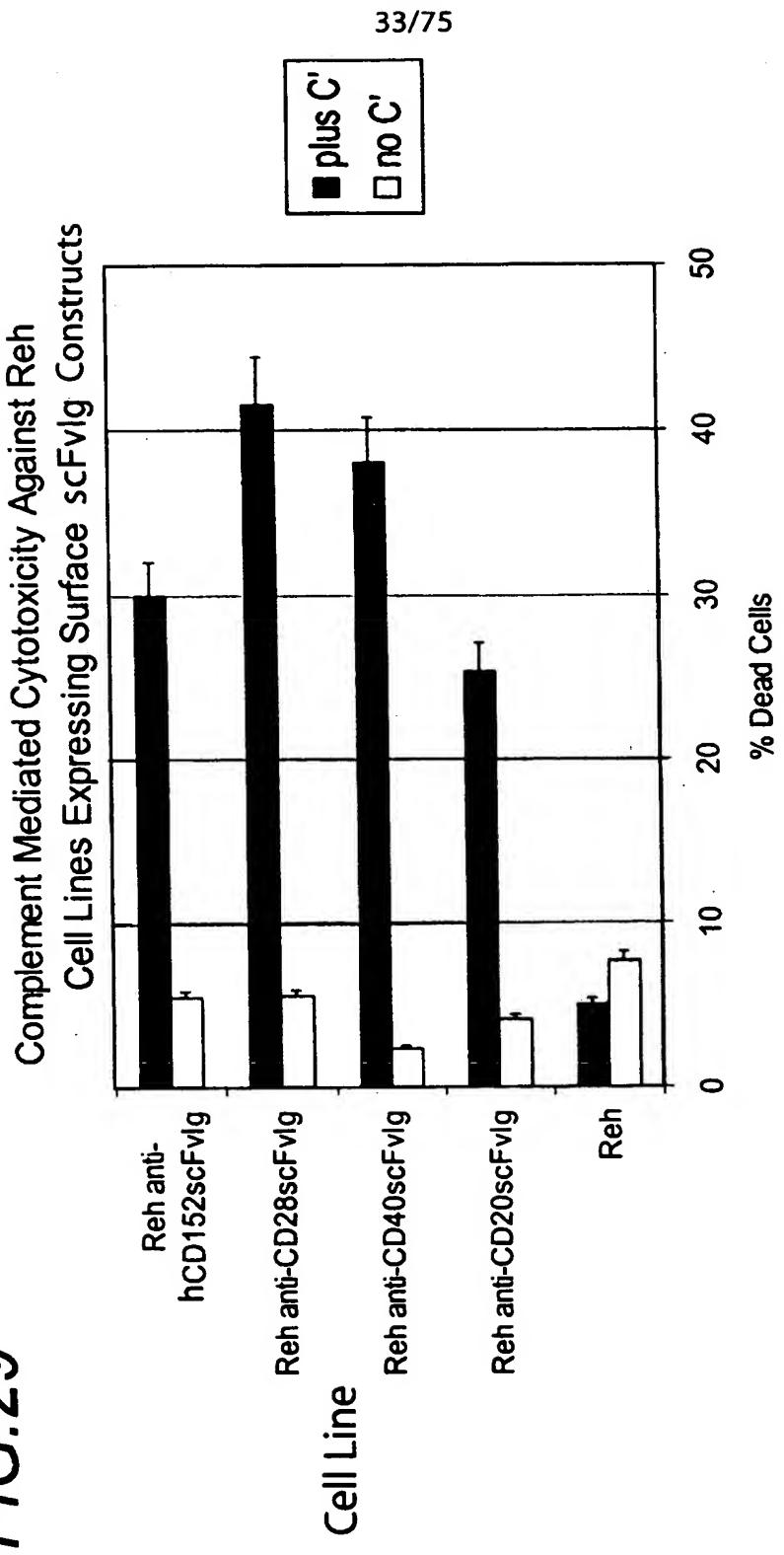


FIG. 30

ADCC Activity of Cell Surface Expressed ScFvIg Constructs

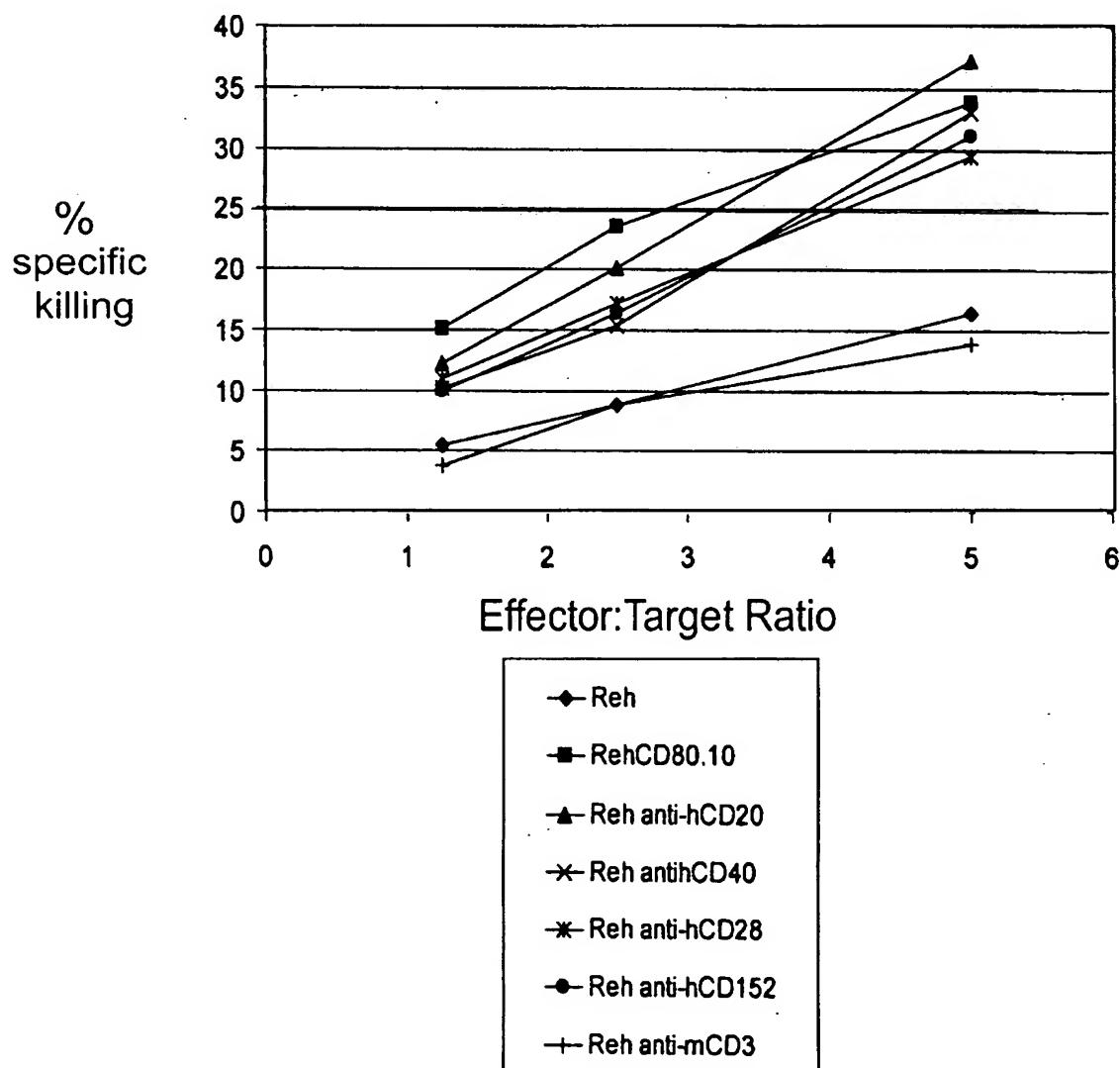


FIG. 31

Ig Constructs and Nomenclature:

35/75

Name Identifier	Hinge Sequence	CH2 Sequence	CH3 Sequence
(CCC-P)WH WCH2 WCH3	IgG1 WT Hinge (CCC)	Wild Type CH2	Wild Type CH3
(SSS-SH) WCH2 WCH3	IgG1 Mutant Hinge (SSS)	Wild type CH2 (IgG1)	Wild type CH3 (IgG1)
VHL11S (SSS-SH) WCH2 WCH3	IgG1 Mutant Hinge (SSS)	Wild type CH2 (IgG1)	Wild type CH3 (IgG1)
(SSC-P)H WCH2 WCH3	IgG1 Mutant Hinge (SSC)	Wild type CH2 (IgG1)	Wild type CH3 (IgG1)
(SCS-SH) WCH2 WCH3	IgG1 Mutant Hinge (SCS)	Wild type CH2 (IgG1)	Wild type CH3 (IgG1)
(CSS-SH) WCH2 WCH3	IgG1 Mutant Hinge (CSS)	Wild type CH2 (IgG1)	Wild type CH3 (IgG1)
(SSS-SH) P238SCH2WCH3	IgG1 Mutant Hinge (SSS)	Mutant CH2 (IgG1) Pro→Ser 238	Wild type CH3 (IgG1)
IgAH WCH2 WCH3	IgA Hinge	Wild type CH2 (IgG1)	Wild type CH3 (IgG1)
IgAH IgA CH2CH3	IgA Hinge	Wild type CH2 (IgA)	Wild type CH3 (IgA)
IgAH IgA CH2T4CH3	IgA Hinge	Wild type CH2 (IgA)	Truncated CH3 (IgA) Missing 4 aa at COOH

FIG. 32

CDC Activity of CTLA4Ig Constructs against Reh and Reh CD80 Transfected Cells

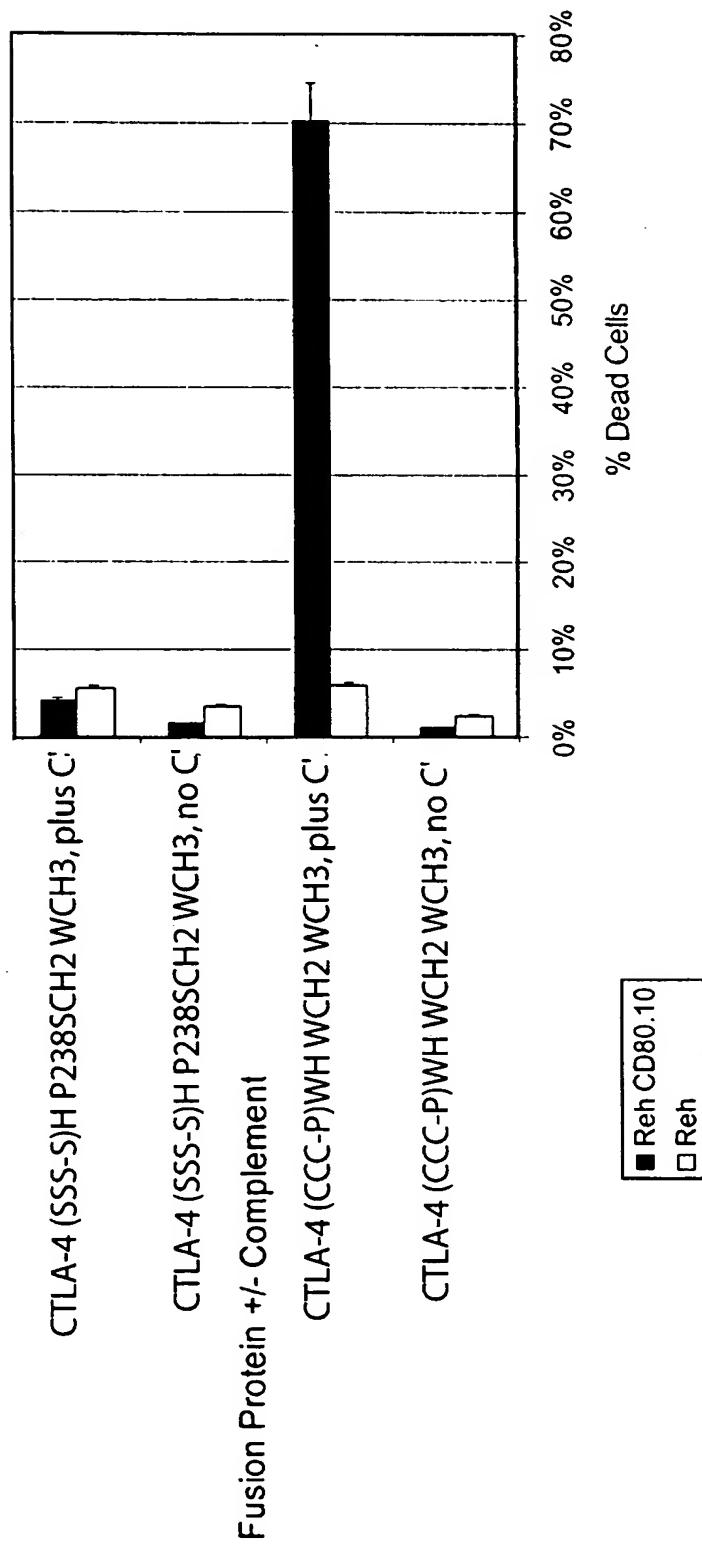


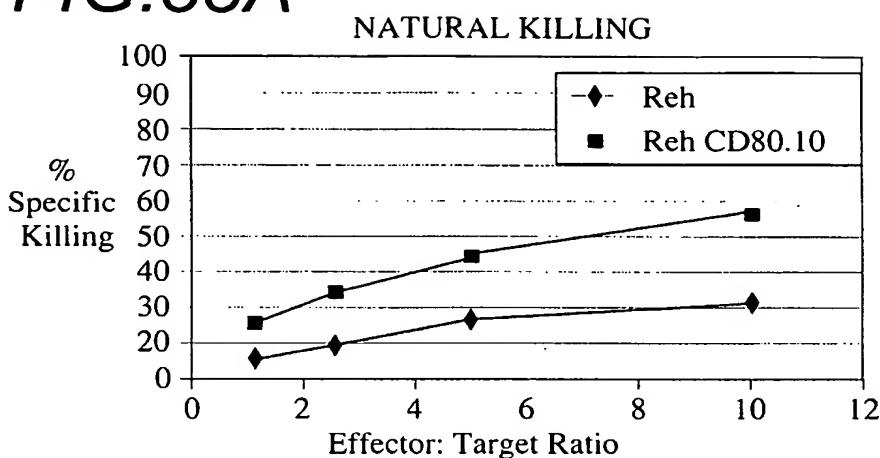
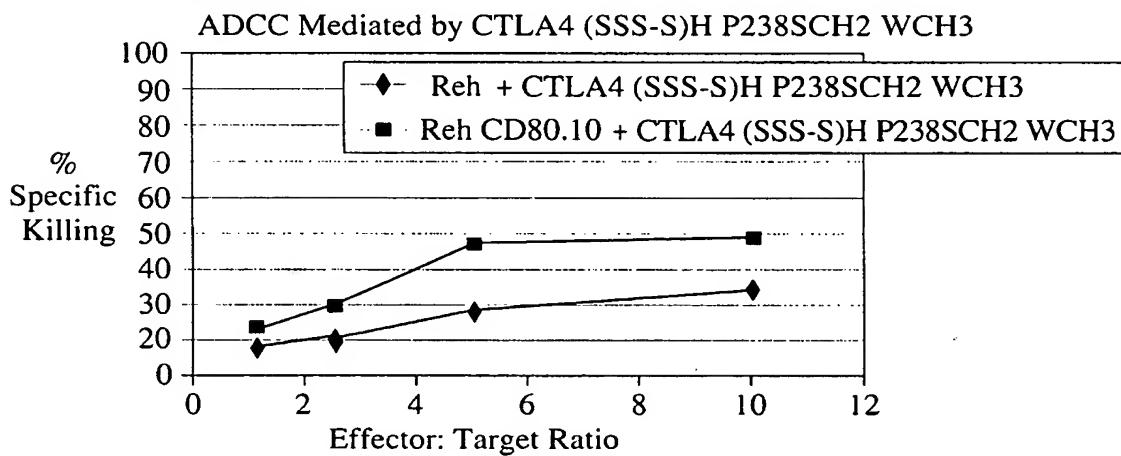
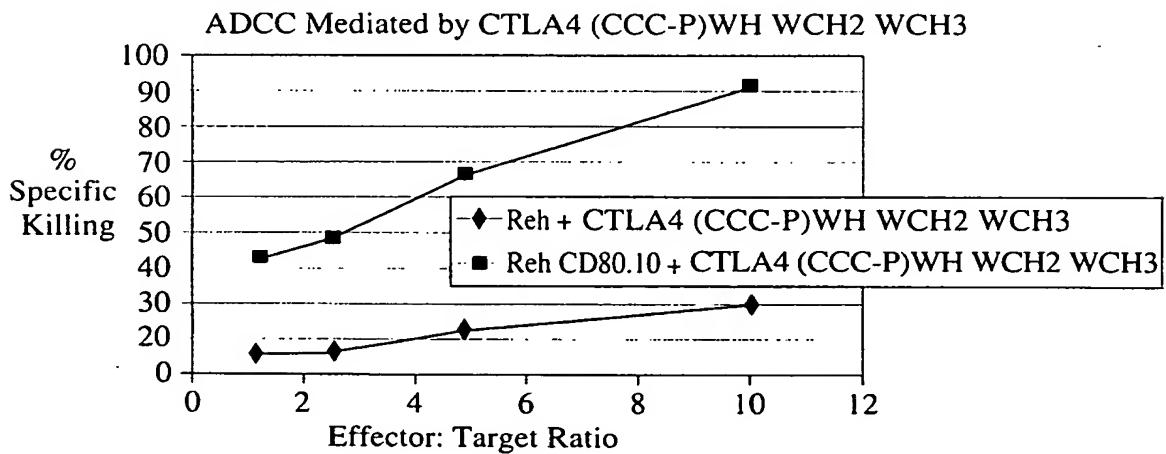
FIG.33A**FIG.33B****FIG.33C**

FIG. 34

Binding of Constructs Alternative to CD2 CHO Cells

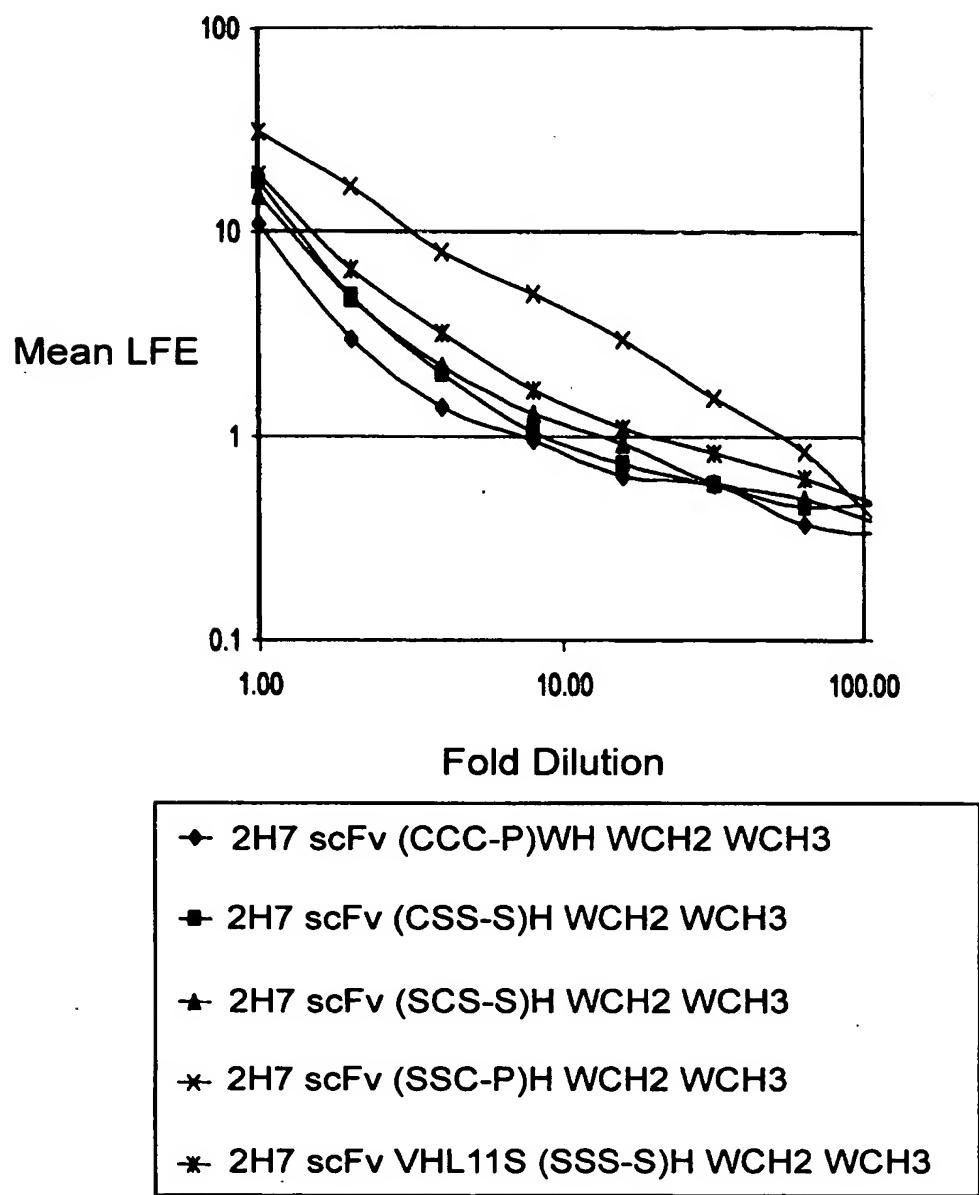


FIG.35

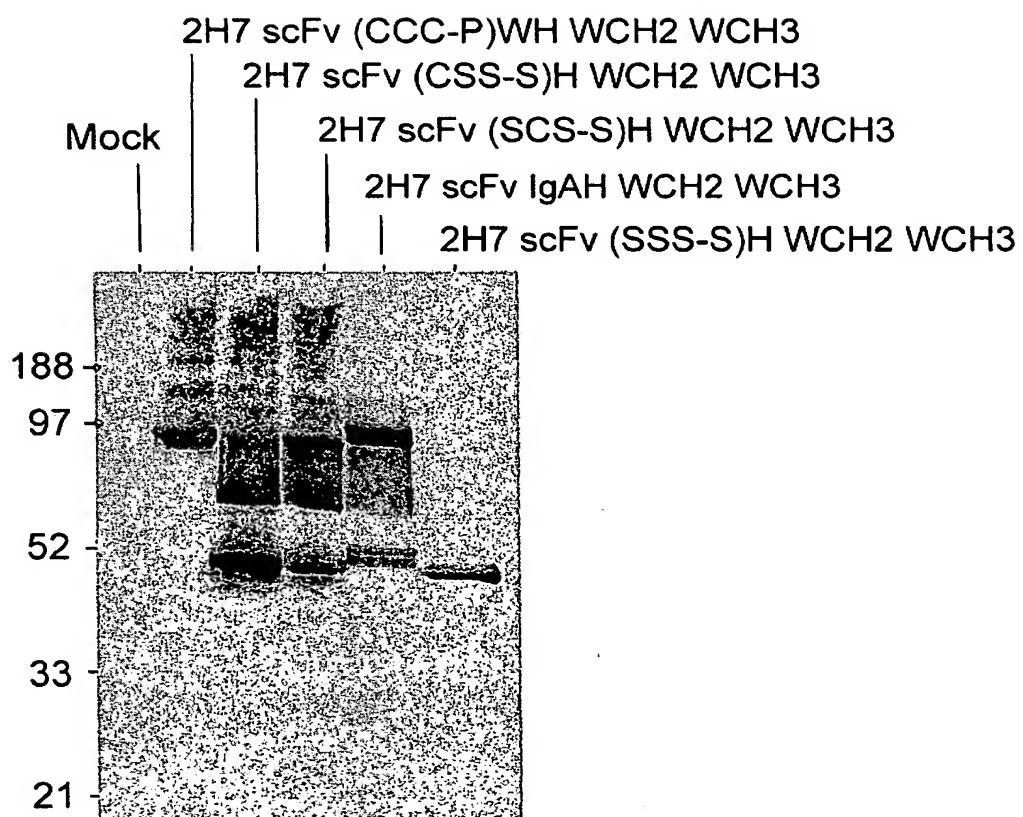


FIG. 36

Binding to CD20 CHO cells by Constructs
that link anti-CD20 scFv to IgA Fc Domains

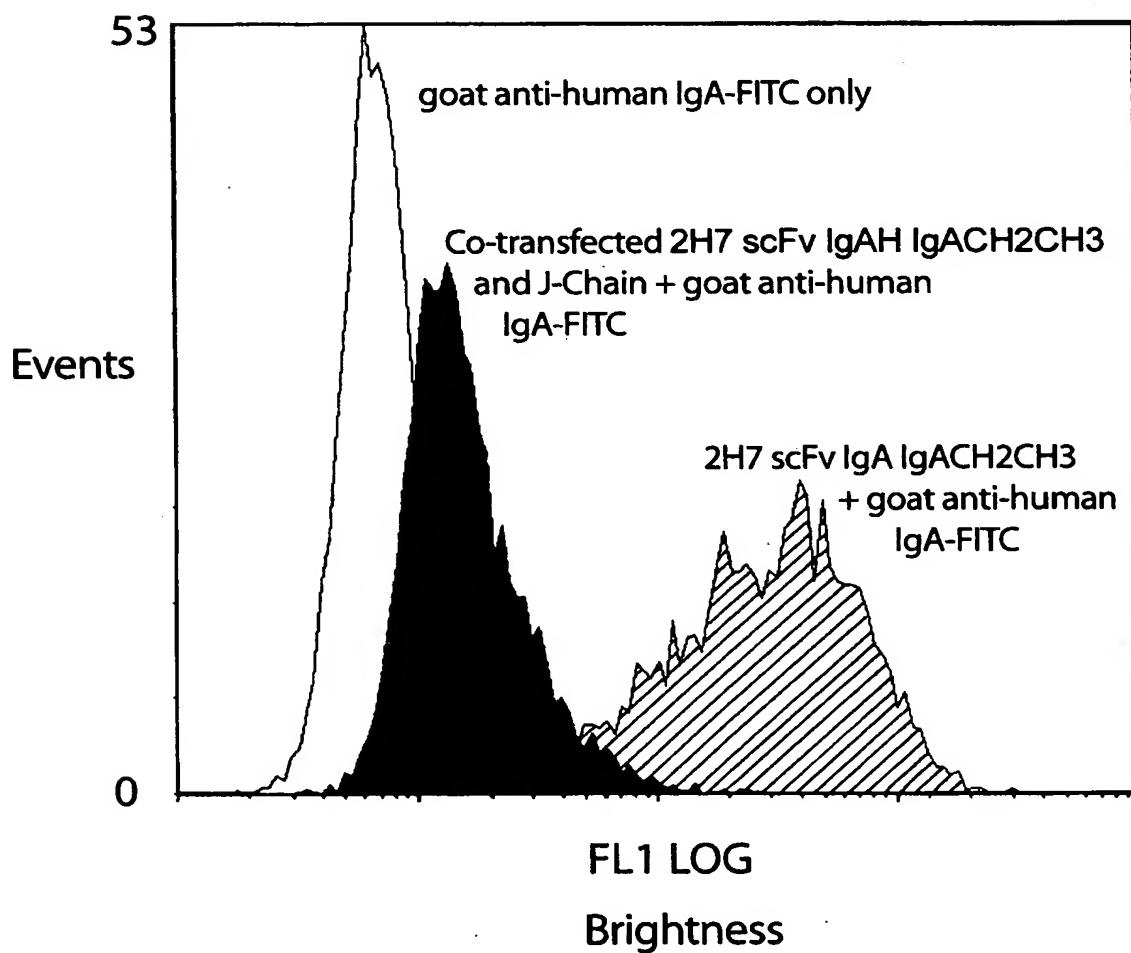


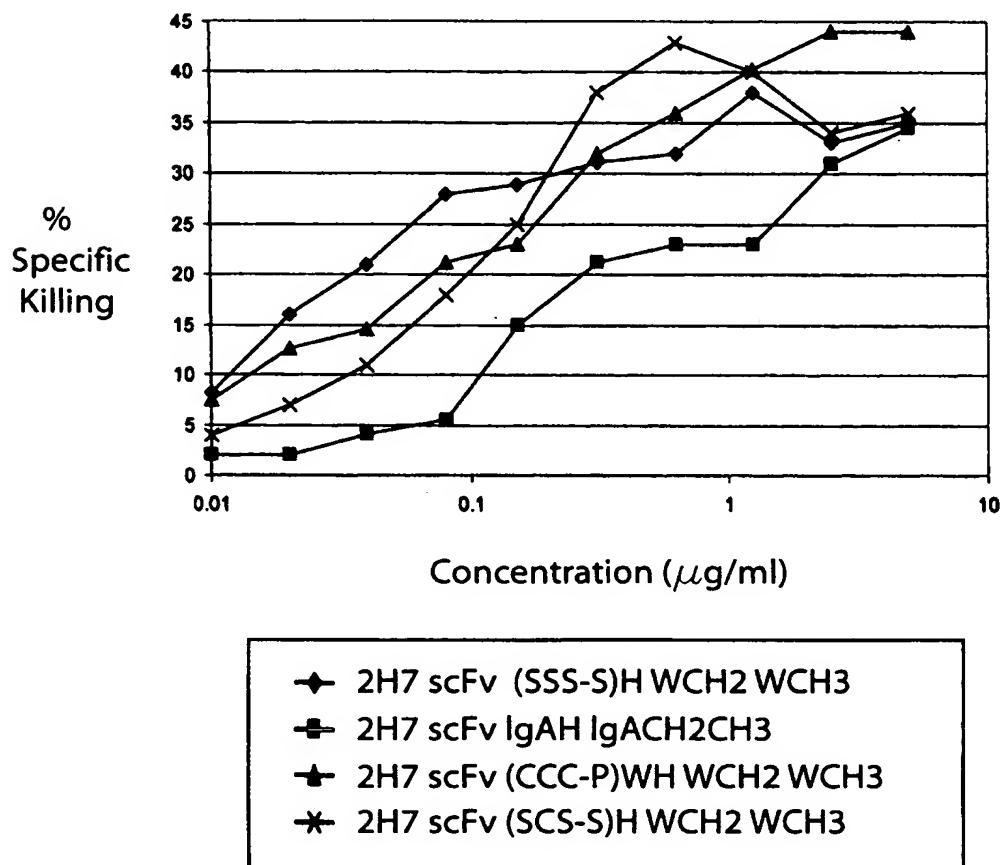
FIG.37**Titration of CD20 Specific scFvIg Constructs for ADCC Activity Using Whole Blood Effectors**

FIG.38

ADCC ASSAY OF ANTI-CD20 CONSTRUCTS WITH DIFFERENT TAILS
 (WHOLE BLOOD EFFECTORS/BJAB TARGETS)

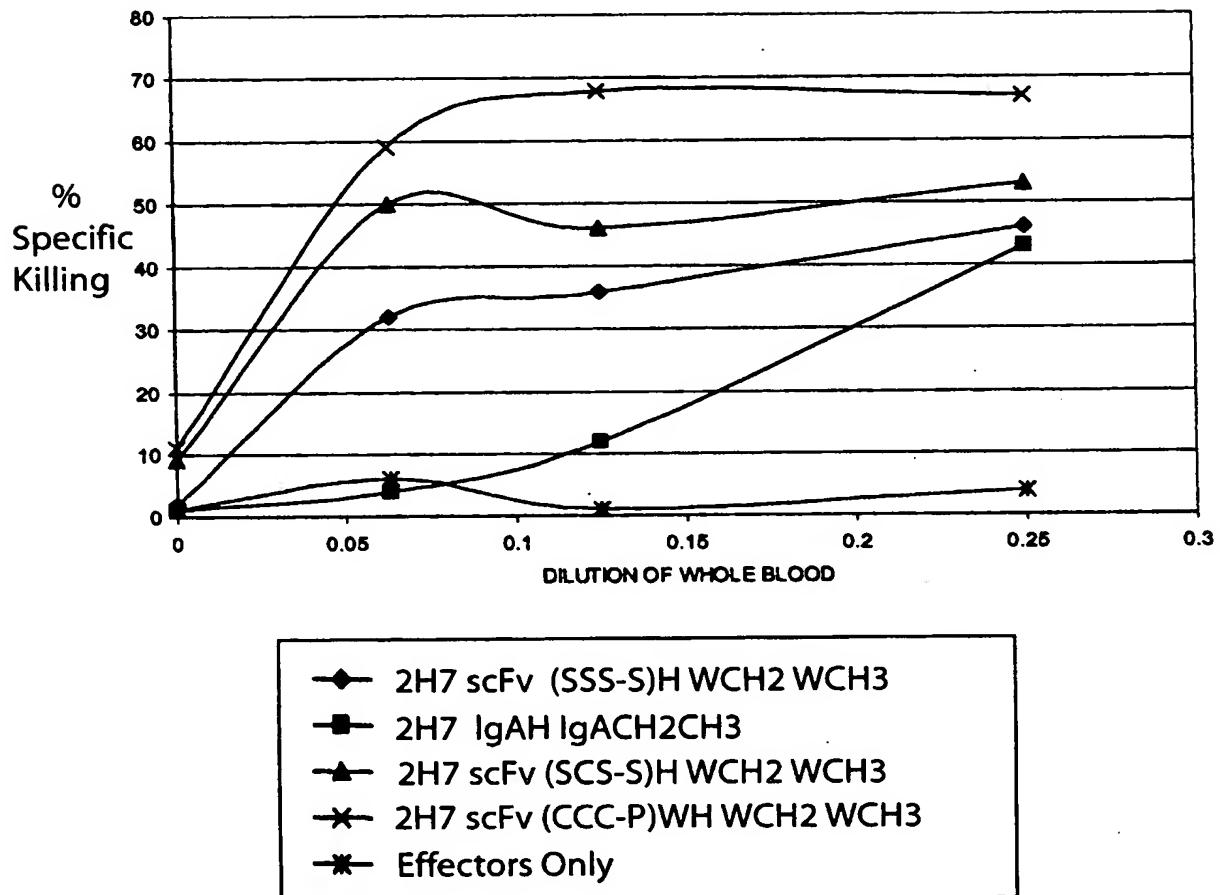
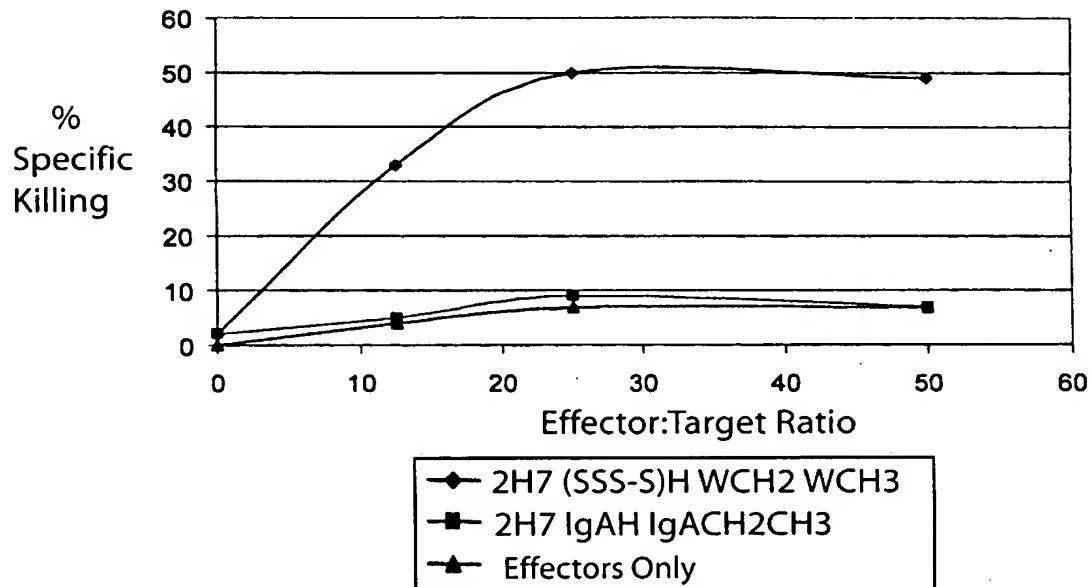


FIG. 39A

**ADCC ASSAY OF ANTI-CD20 CONSTRUCTS
(PBMC EFFECTORS/BJAB TARGETS)**

**FIG. 39B**

**ADCC ASSAY OF ANTI-CD20 CONSTRUCTS
(WHOLE BLOOD EFFECTORS/BJAB TARGETS)**

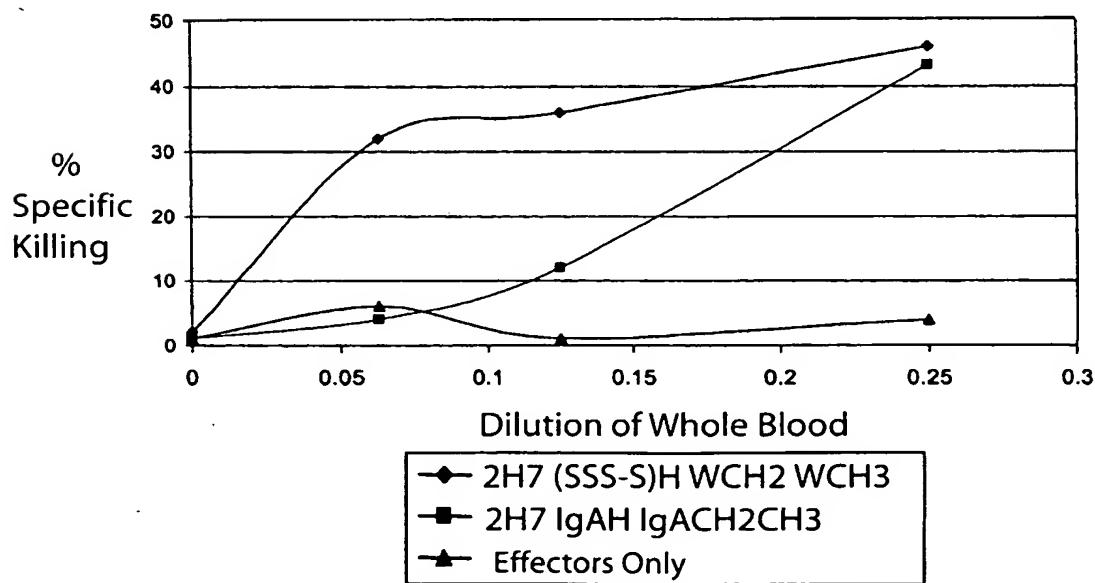
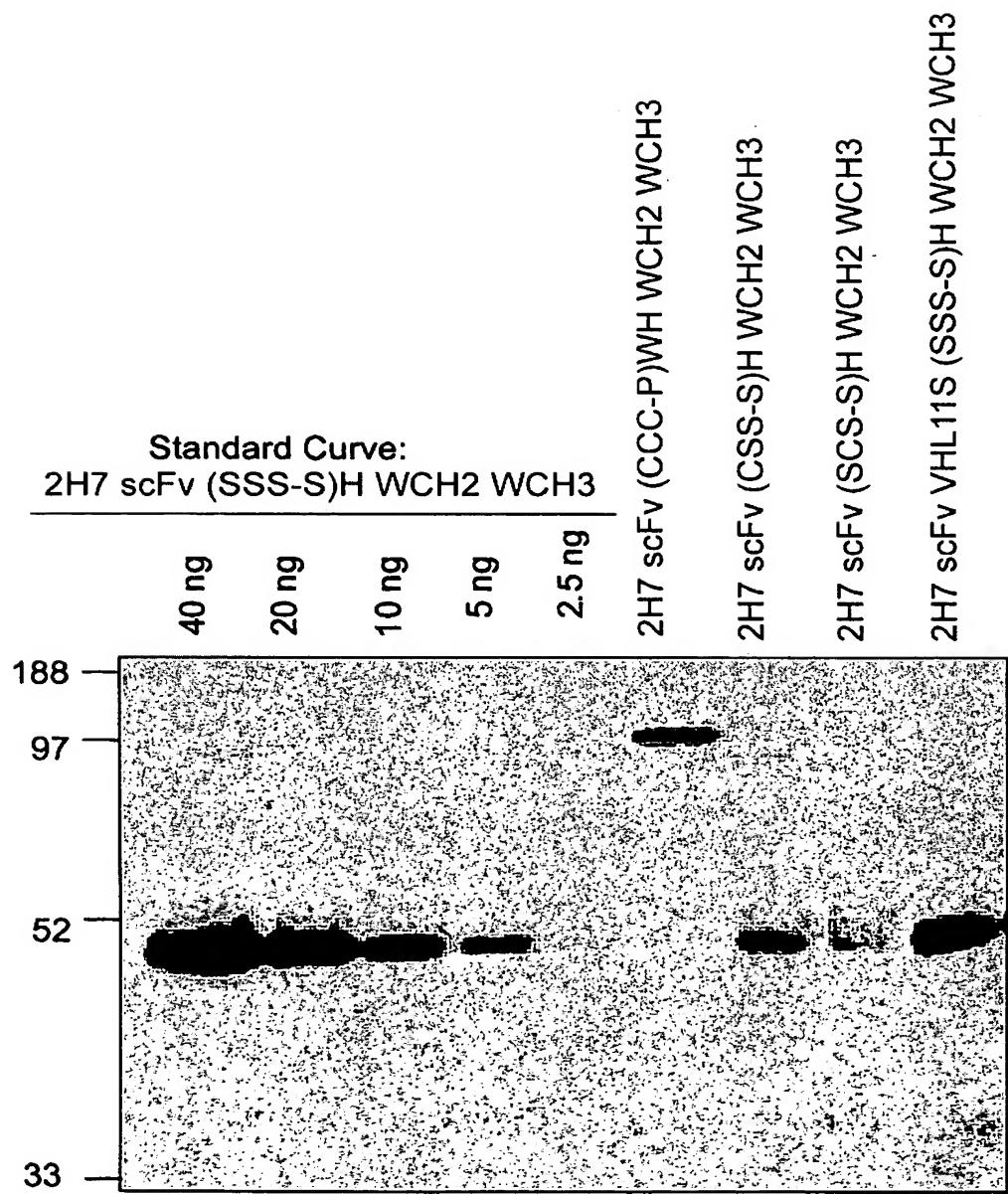


FIG. 40

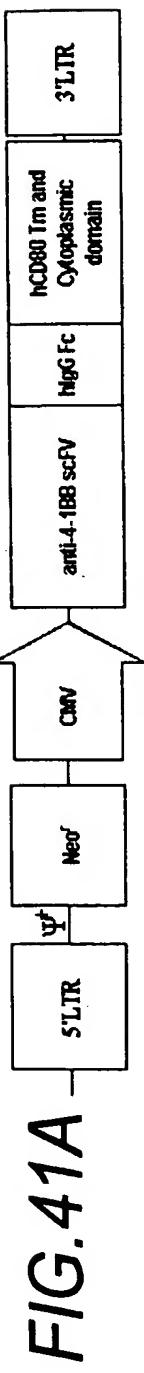
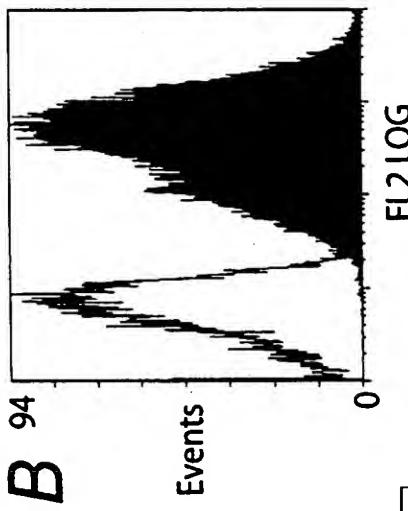
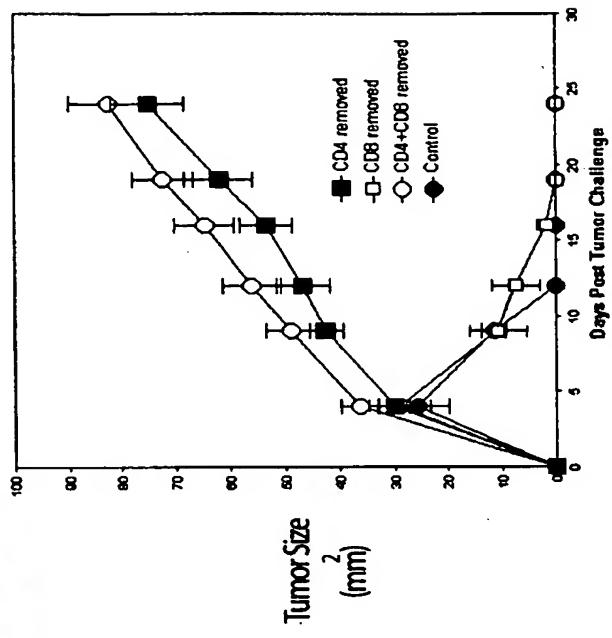
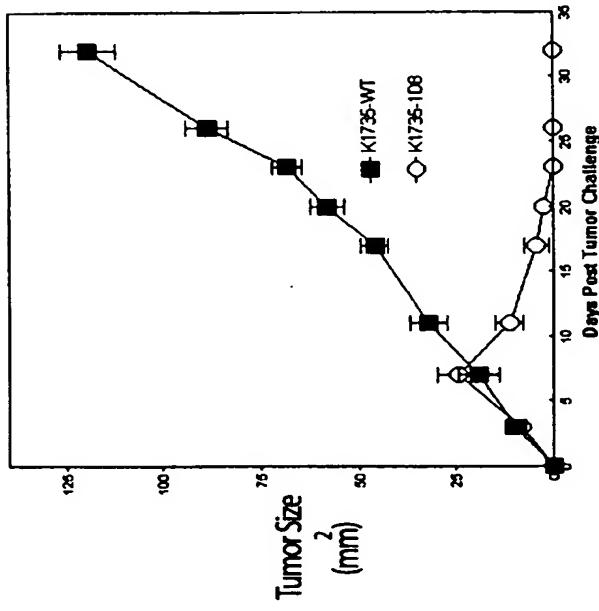
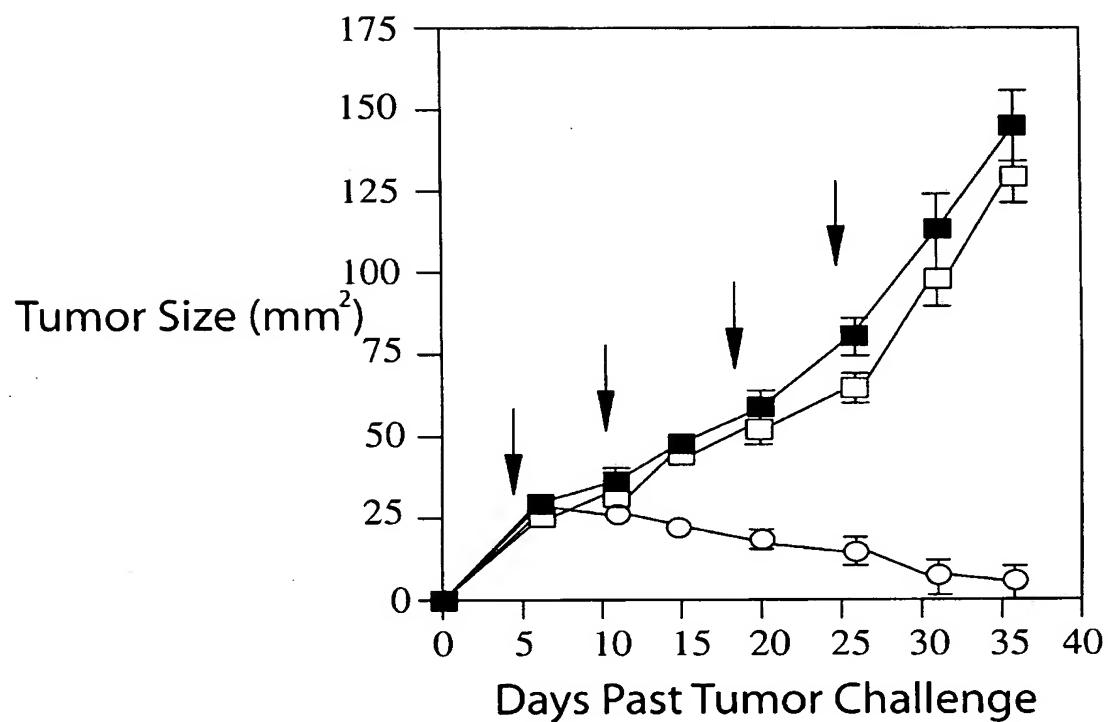
**FIG. 41B****FIG. 41D****FIG. 41C**

FIG.42

47/75

FIG. 43

Mixtures of K1735-WT and K1735-1D8 transfected tumor lines inhibit tumor outgrowth in C3H mice

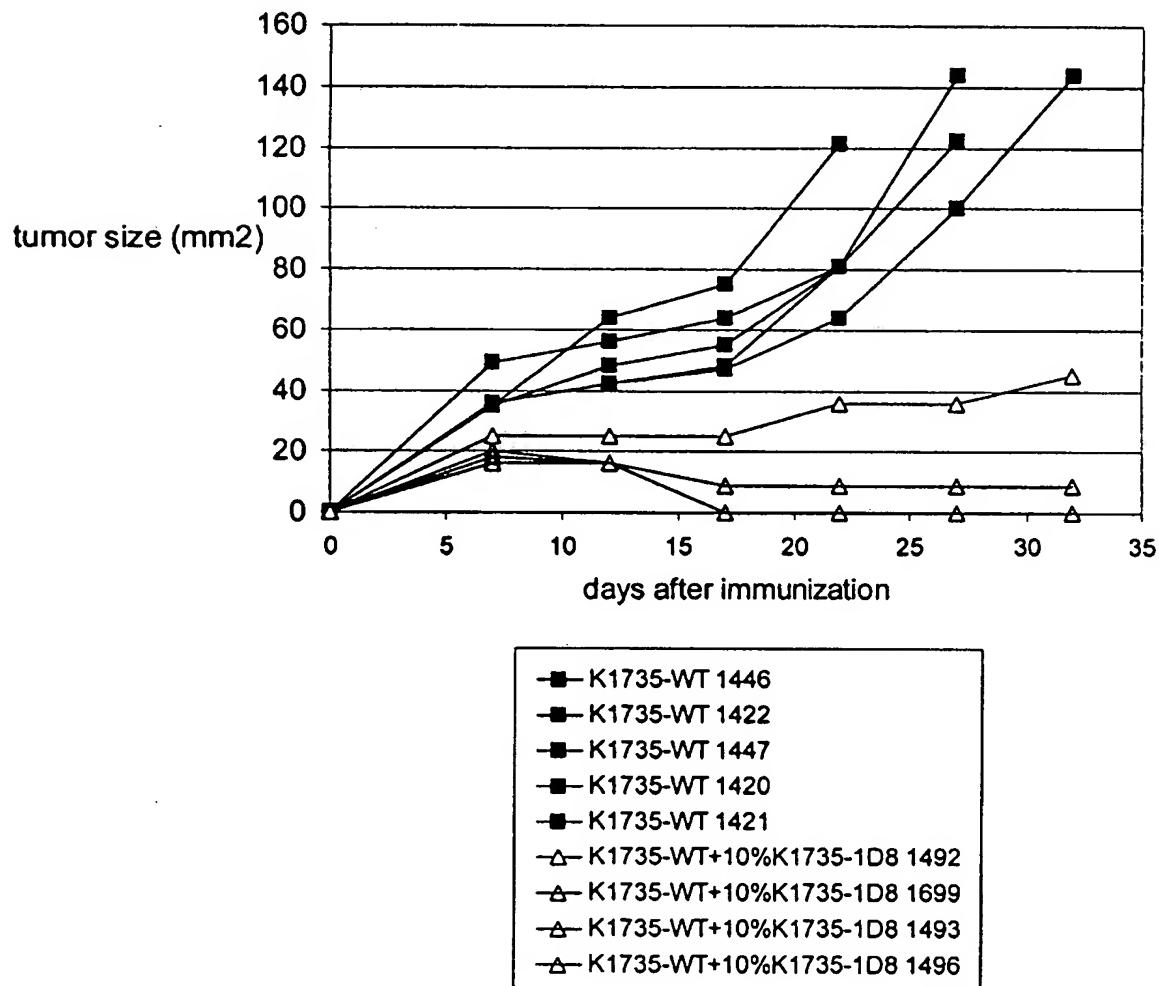


FIG. 44

Expression of 1D8 scFv (SSS-S)H P238SCH2 WCH3 (Anti-CD37)
on the Surface of Panned Ag104 Transfected Tumor Cells

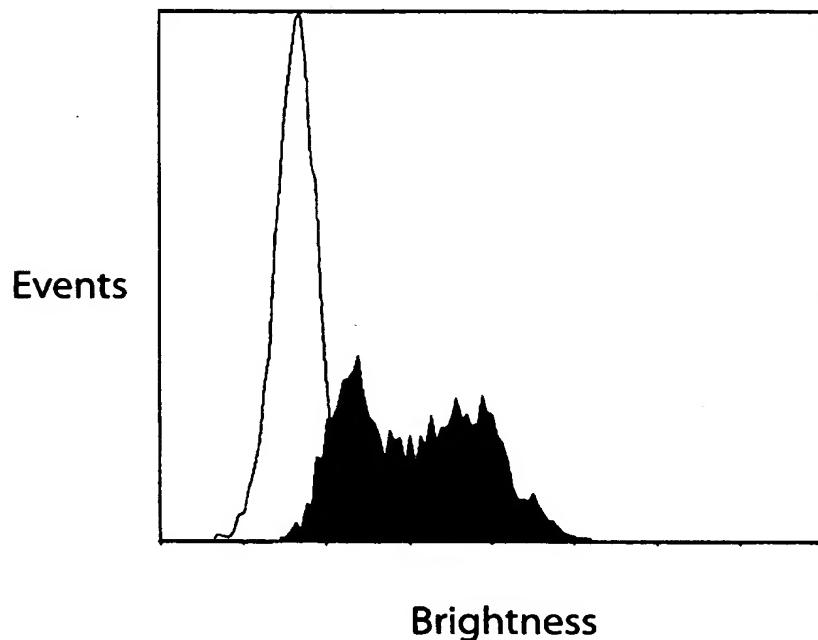


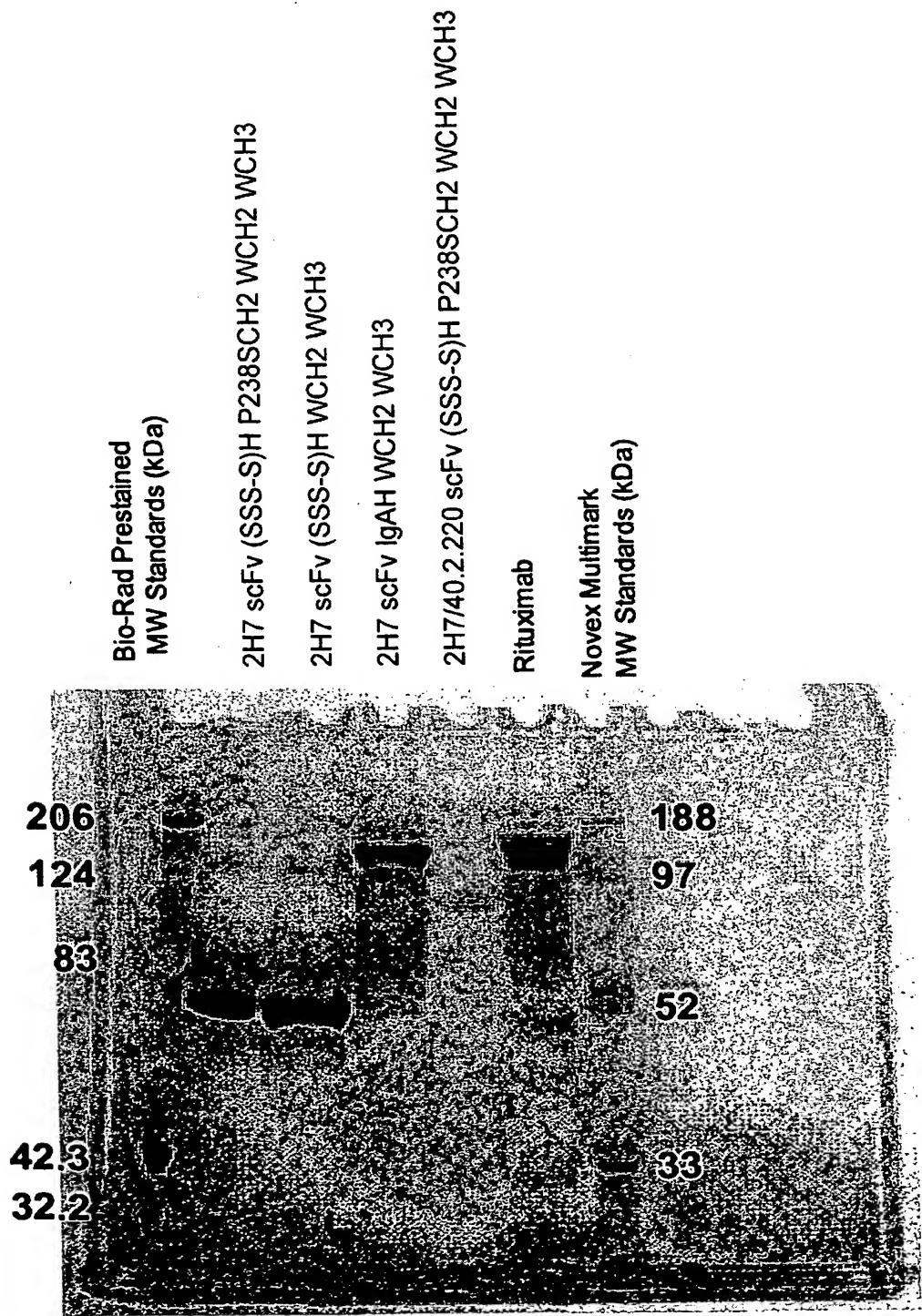
FIG.45

FIG.46

ADCC mediated by 2H7 scFvIg constructs
by human PBMC effector cells against Bjab targets

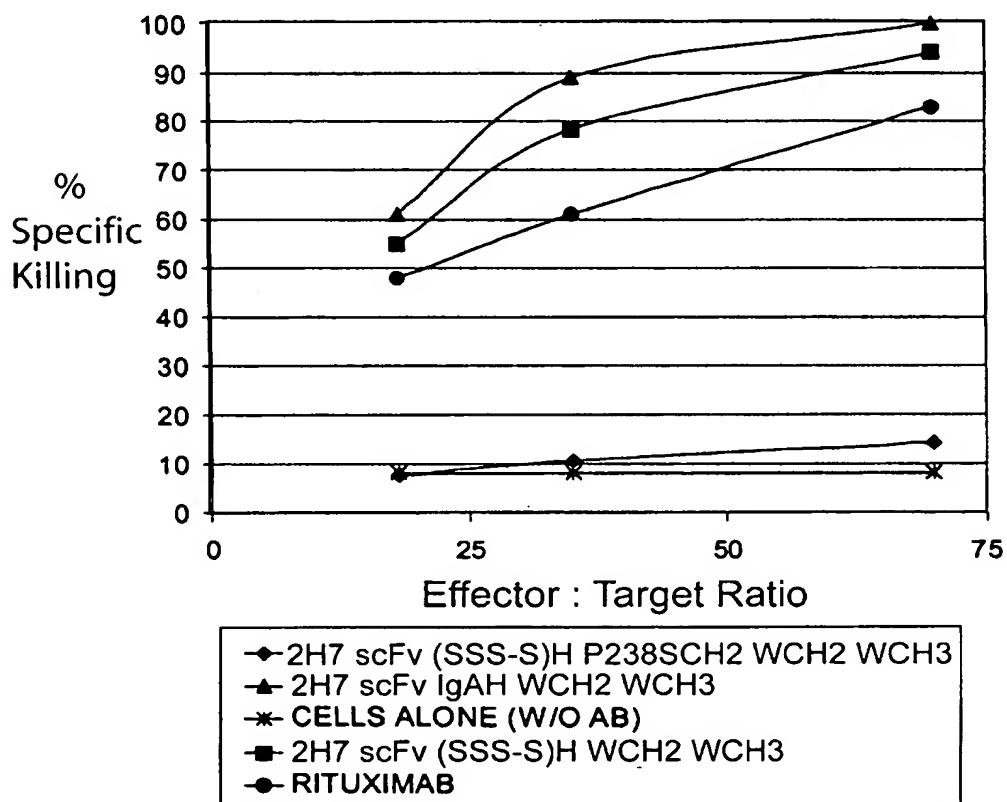


FIG.47

Cell surface expression of
G19-4scFv (SSS-S)H P238SCH2 WCH3
fusion protein on Reh and T51 Cells.

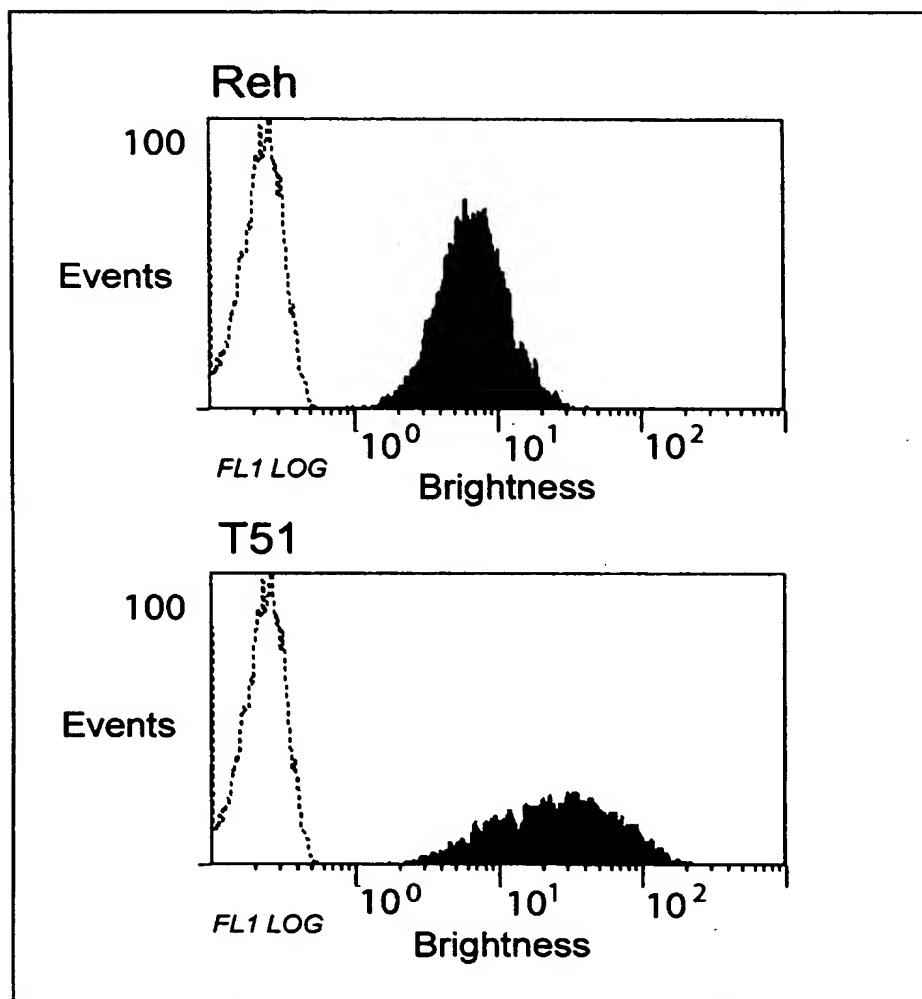


FIG.48

**Targeting of Cytotoxicity to Transfected Cell Lines
by Surface expression of
G19-4 scFv (SSS-S)H P238SCH2 WCH3**

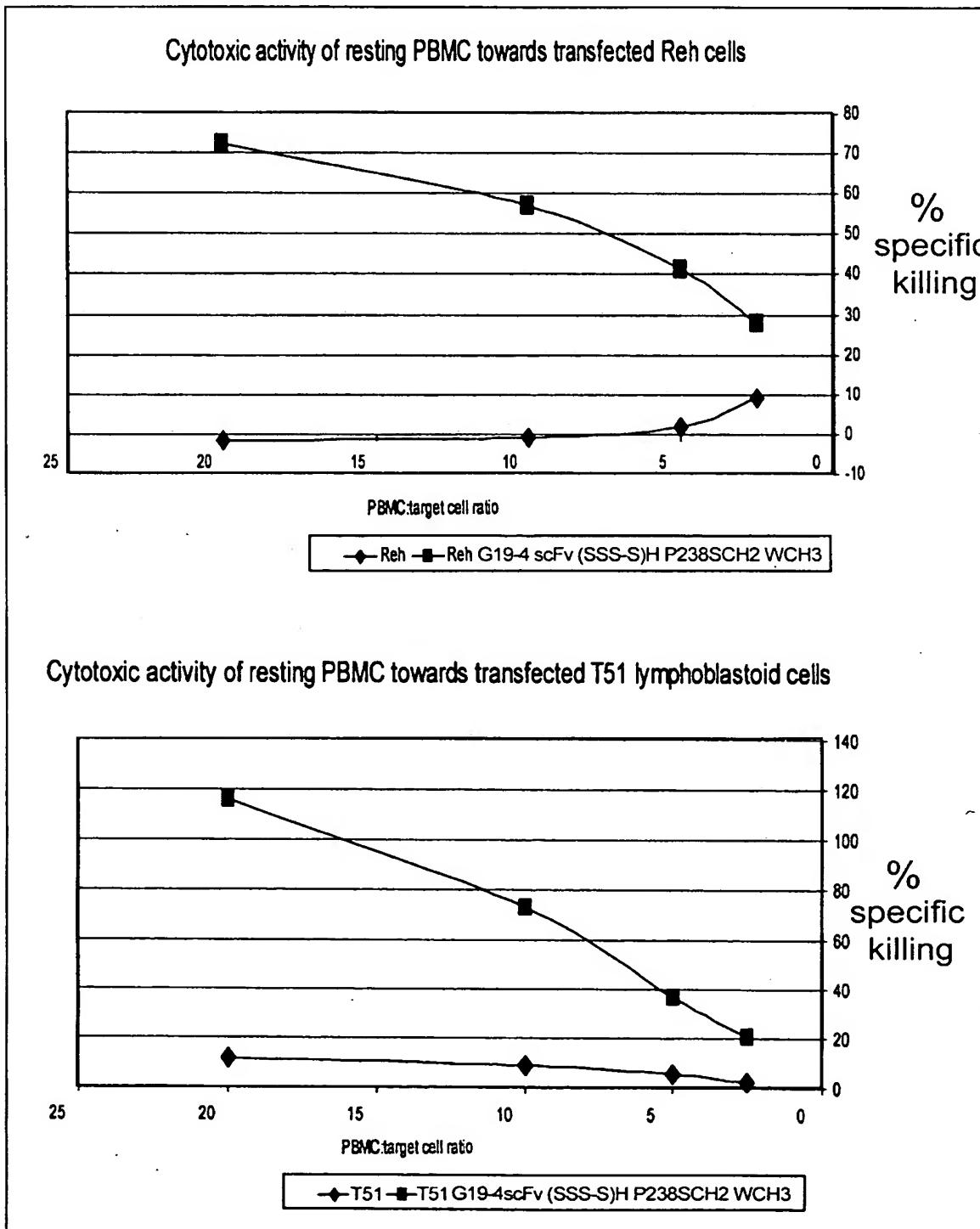


FIG.49

Binding of 5B9 scFv (SSS-S)H WCH2 WCH3,
a mouse anti-human CD137 to stimulated human PBMC

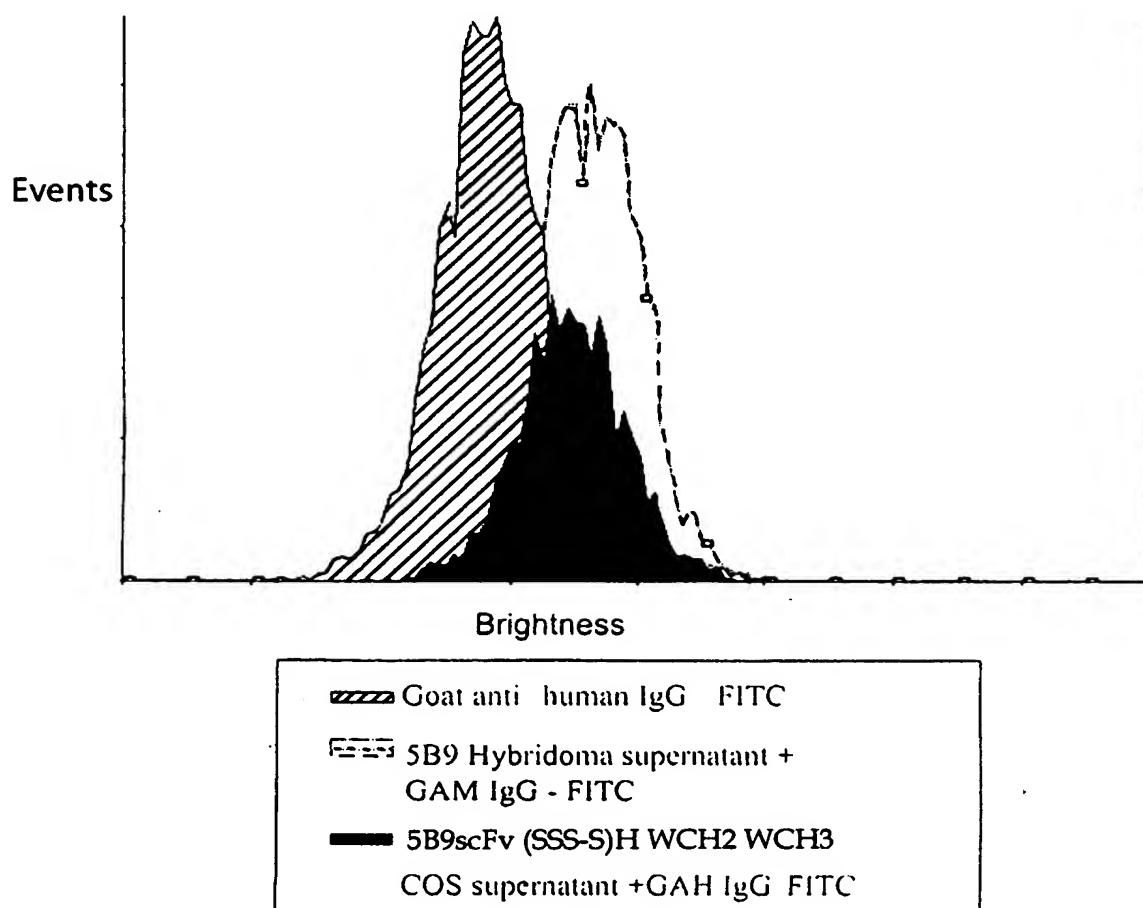
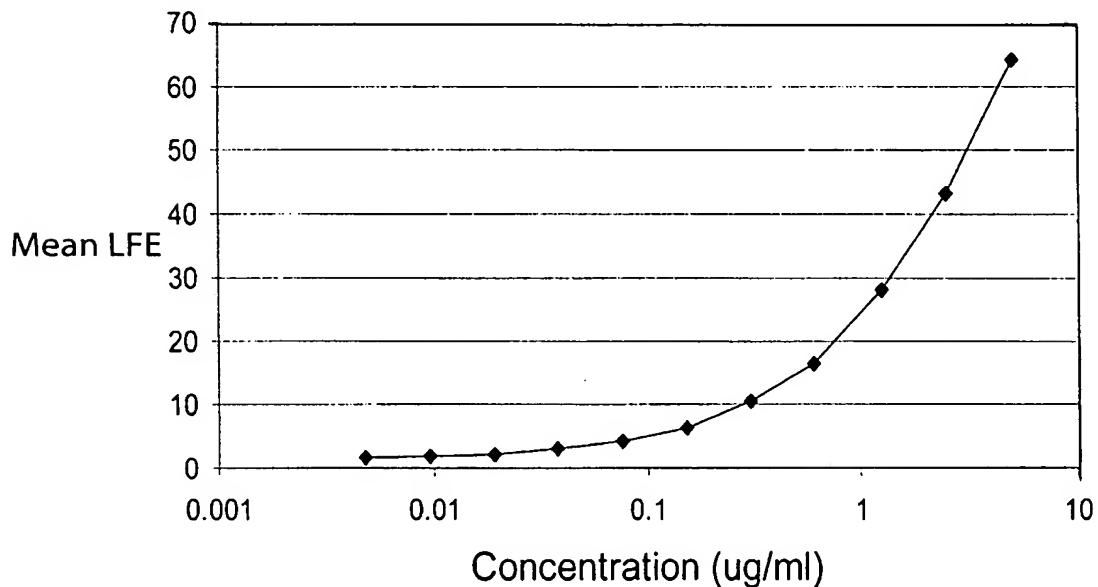


FIG.50A

Effect of V_H L11S Mutation on
2H7 scFv (SSS-S)H WCH2 WCH3 Protein Expression
Standard Curve: 2H7scFv VHL11S (SSS-S)H WCH2 WCH3

**FIG.50B**

Effect of V_H L11S Mutation on
2H7 scFv (SSS-S)H WCH2 WCH3 Protein Expression

CHO supernatant Brightness and Estimation of
Protein concentrations from Standard Curve:

	CHO clone name				
	<u>4F2</u>	4F5	3E5	6B11A	2B8A
Mean LFE					
1/100	71.7	40.6	31.5	99.7	101.5
1/500	27.1	12.4	11.2	40.8	43
approx conc. μ g/ml	600	225	125	1000	1250

FIG.51

**Production Levels of 2H7scFv VH L11S
(SSS-S)H WCH2 WCH3
From CHO Clone Culture Supernatants**

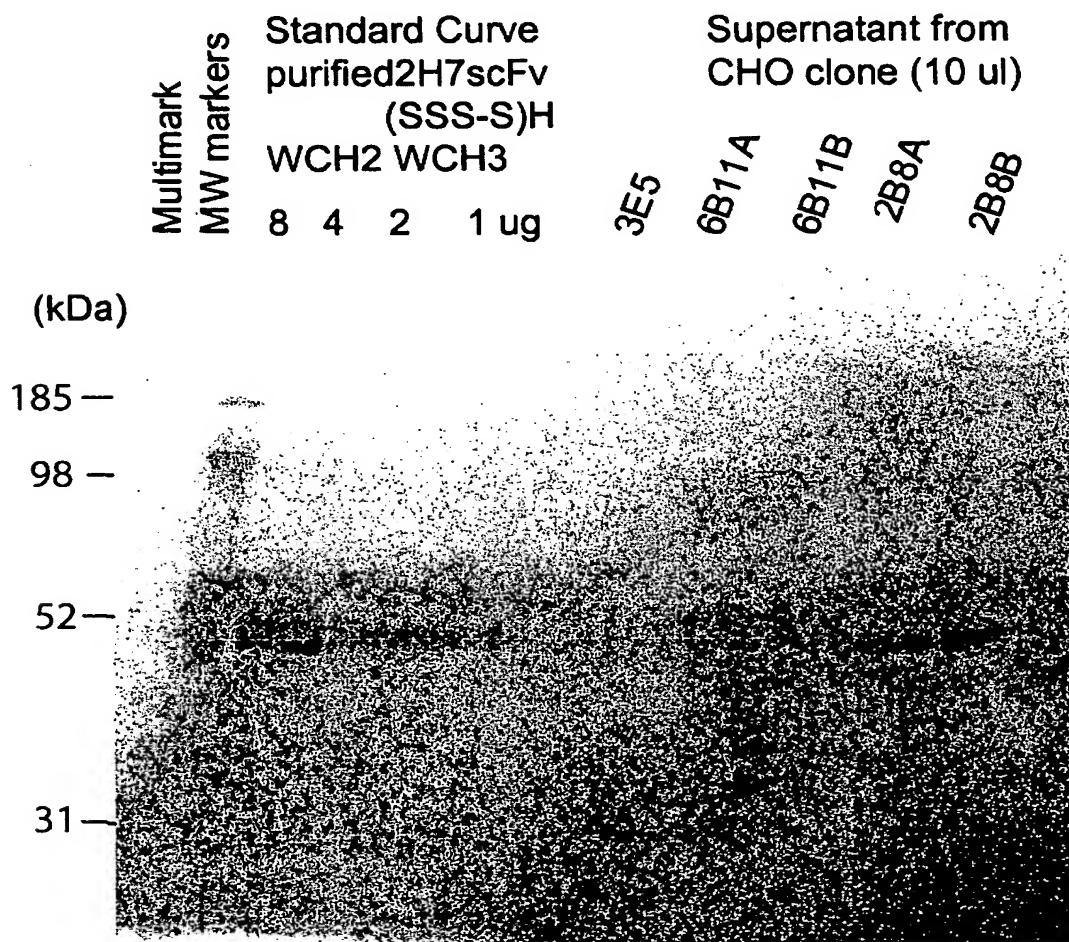


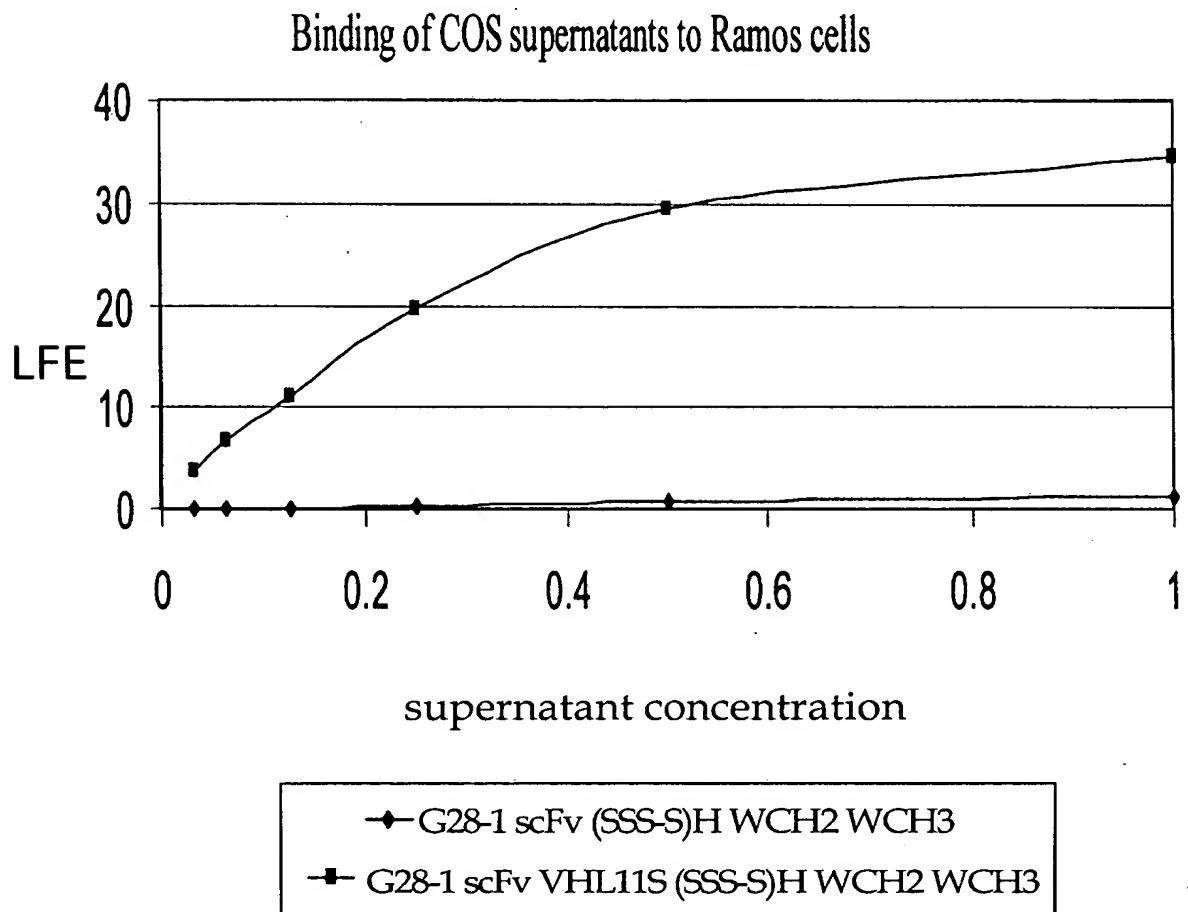
FIG.52**Effect of VHL11S Mutation on G28-1 scFvIg Construct Protein Production from COS cells**

FIG. 53A

Immunoblot of G28-1 scFvIg Constructs

Increased Protein Levels in COS supernatants
transfected with G28-1scFv (SSS-S)H WCH2 WCH3
After Substitution of Leucine with Serine at position 11 of VH (VHL11S)

Purified G28-1	G28-1 scFv (SSS-S)H	G28-1 scFv VHL11S
scFv (SSS-S)H	WCH2 WCH3	WCH2 WCH3
WCH2 WCH3	1 μ l/well	1 μ l/well
80 ng	20 ng	10 ng
40 ng	20 ng	10 ng
20 ng	10 ng	10 ng
10 ng	10 ng	10 ng
A	B	C
B	C	D
C	D	E

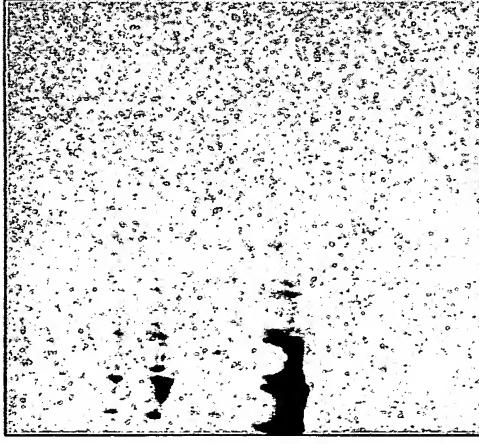
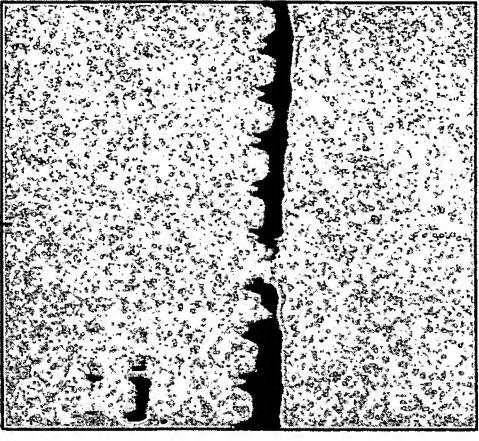


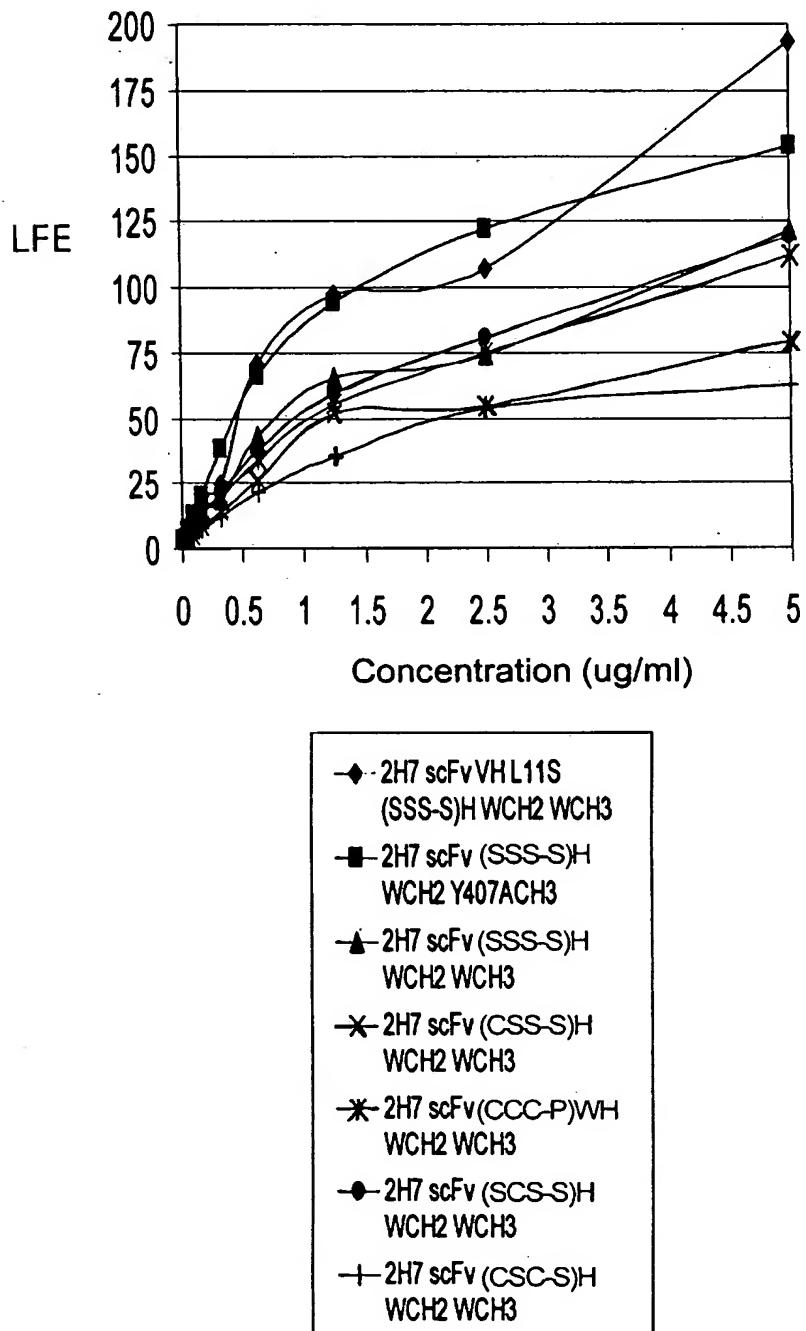
FIG. 53B

Immunoblot of G28-1 scFvIg Constructs

Increased Protein Levels in COS supernatants
transfected with G28-1scFv (SSS-S)H WCH2 WCH3
After Substitution of Leucine with Serine at position 11 of VH (VHL11S)

Purified G28-1	G28-1 scFv VHL11S
scFv (SSS-S)H	WCH2 WCH3
WCH2 WCH3	1 μ l/well
80 ng	20 ng
40 ng	20 ng
20 ng	10 ng
10 ng	10 ng
A	B
B	C
C	D
D	E



Binding of 2H7 scFvIg Constructs with Altered Hinges and CH3 domains to CD20 CHO Cells

ADCC Activity of 2H7 scFvlg constructs Against
BJAB Targets and PBMC Effectors

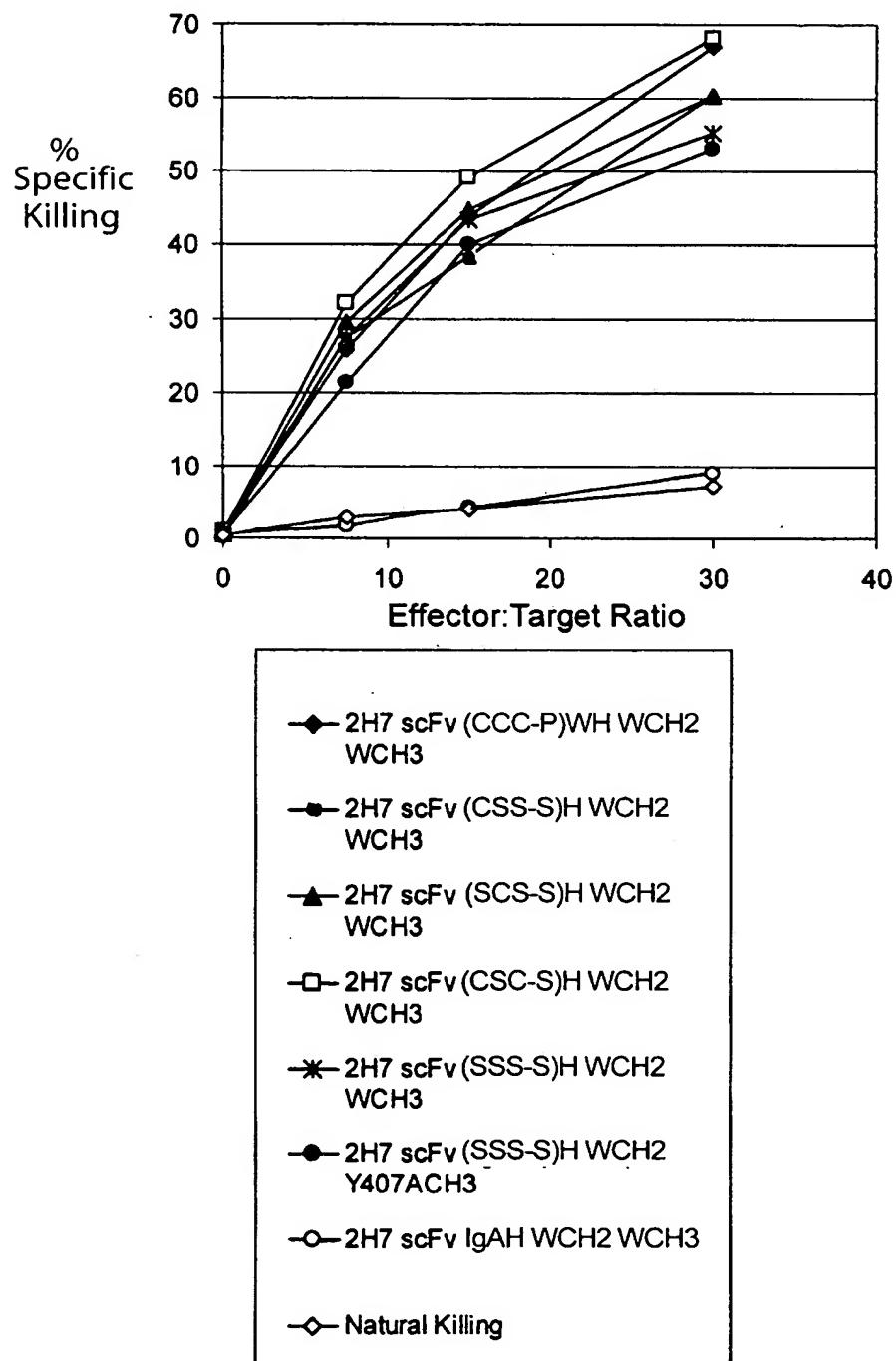


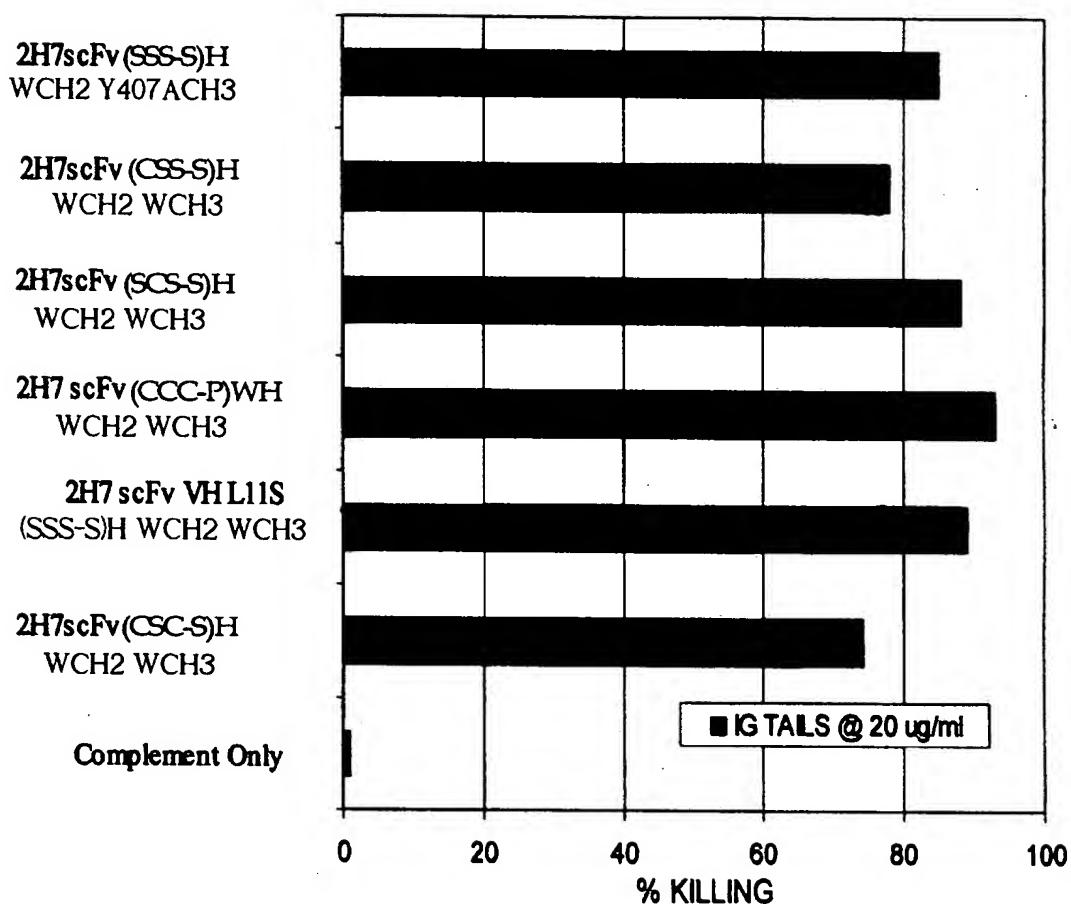
FIG.56**Complement Activity of 2H7 scFvIg Constructs
With Ramos Target Cells**

FIG. 57

Binding of 2H7 scFvIg Derivatives CD20CHO Cells

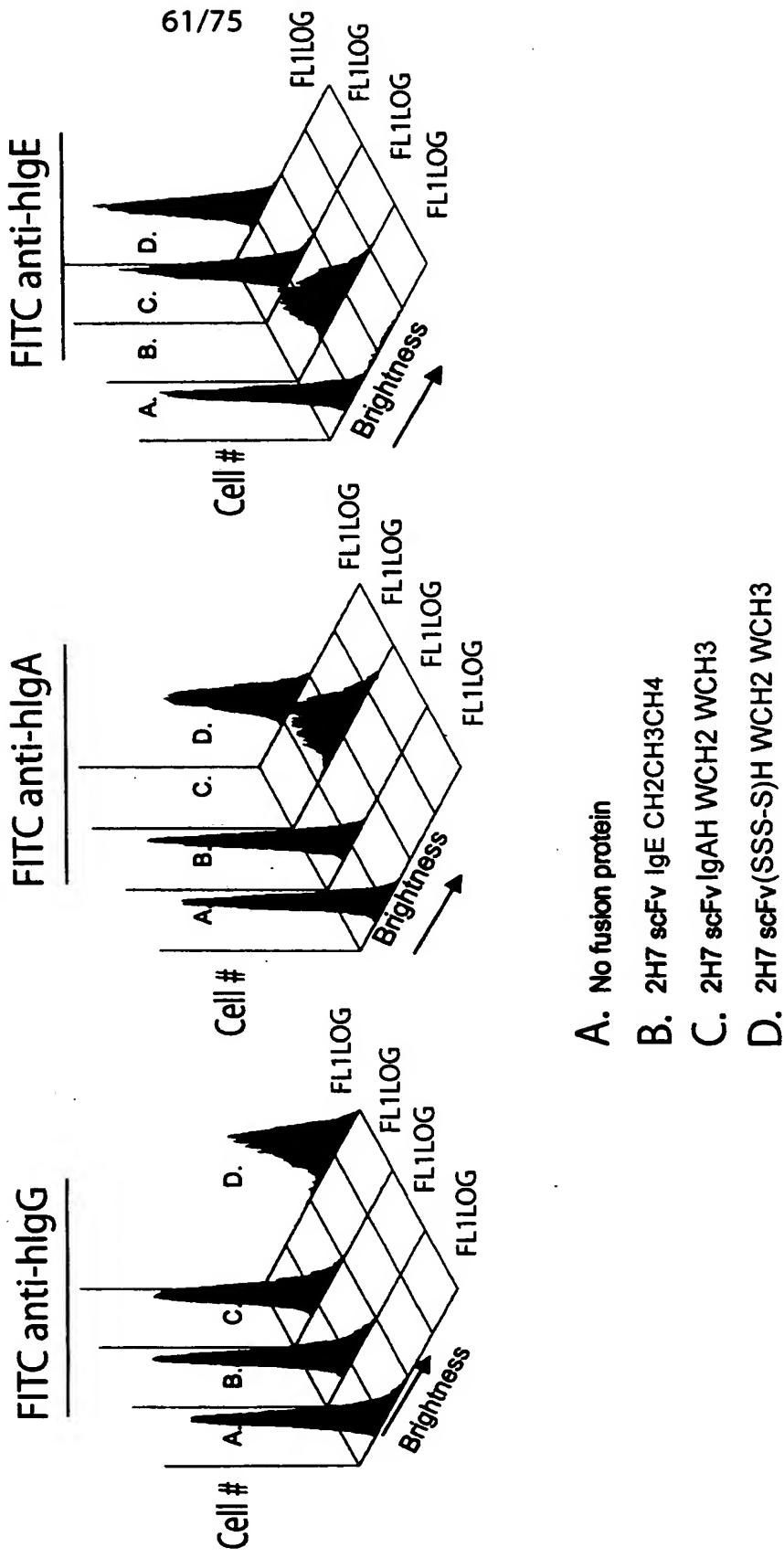


FIG. 58A

62/75

2H7 scFv VH L11S human IgE CH₂CH₃CH₄
Binding to CD20 CHO at 30 ug/ml

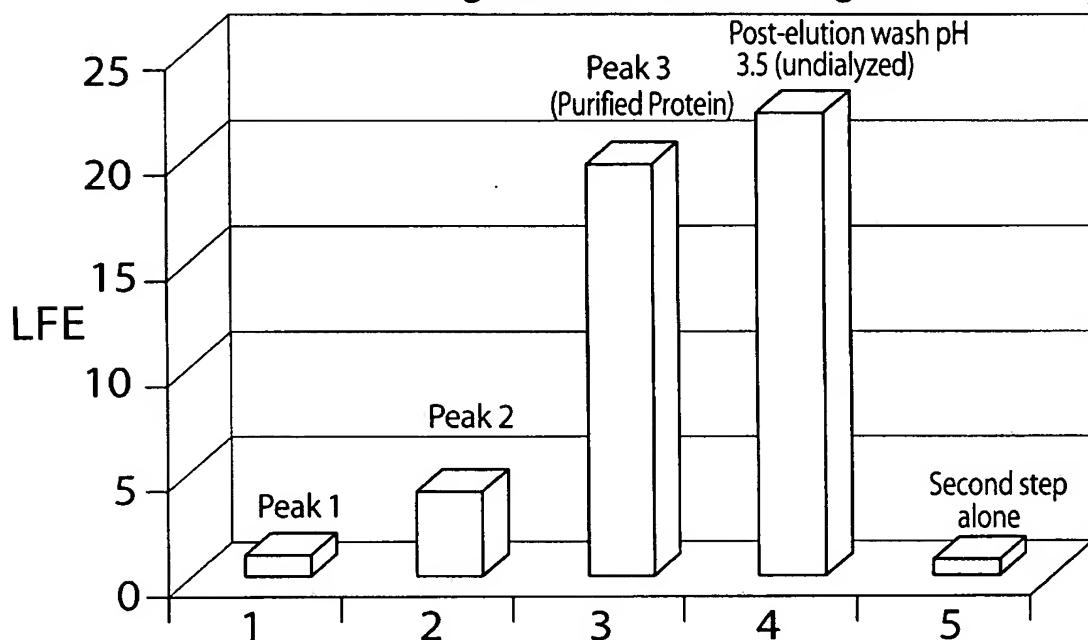


FIG. 58B

ADCC Activity of 2H7 scFv VHL11S IgE CH₂CH₃CH₄
Protein Fractions with PBMC Effectors and Bjab Targets

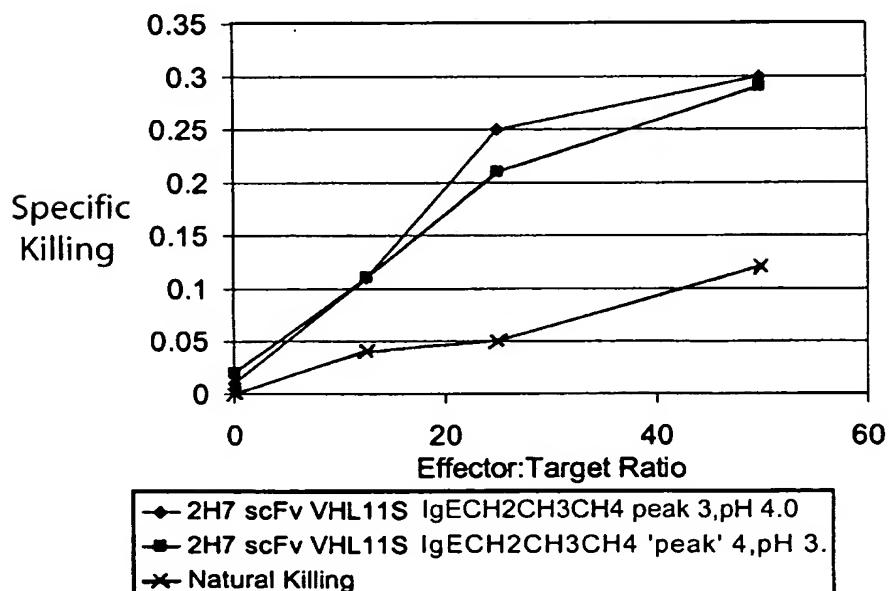


FIG. 59

Binding Data For COS derived 2H7 scFv VHL11S
mIgECH2CH3CH4 and mIgAH WCH2 WCH3

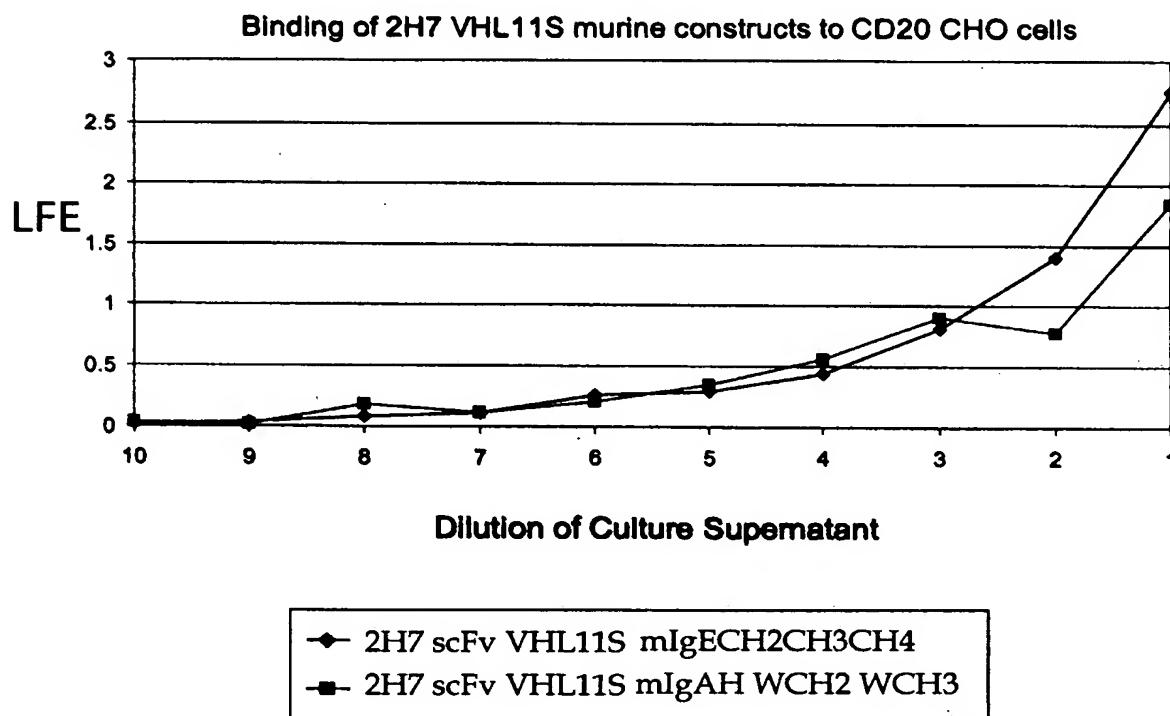


FIG. 60A

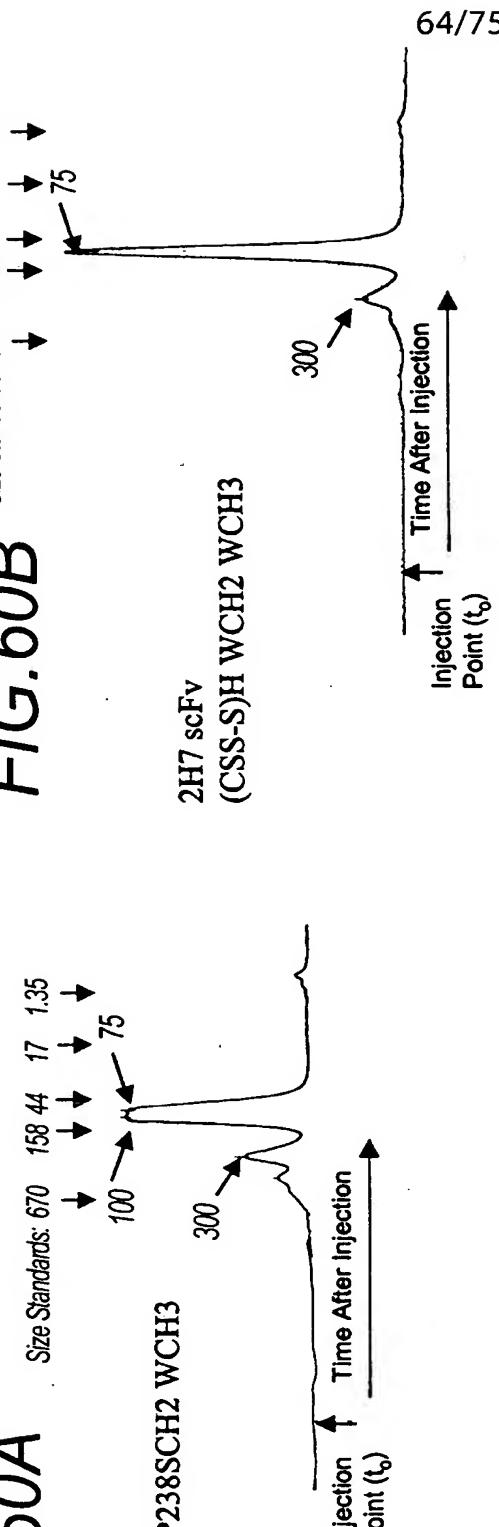


FIG. 60C

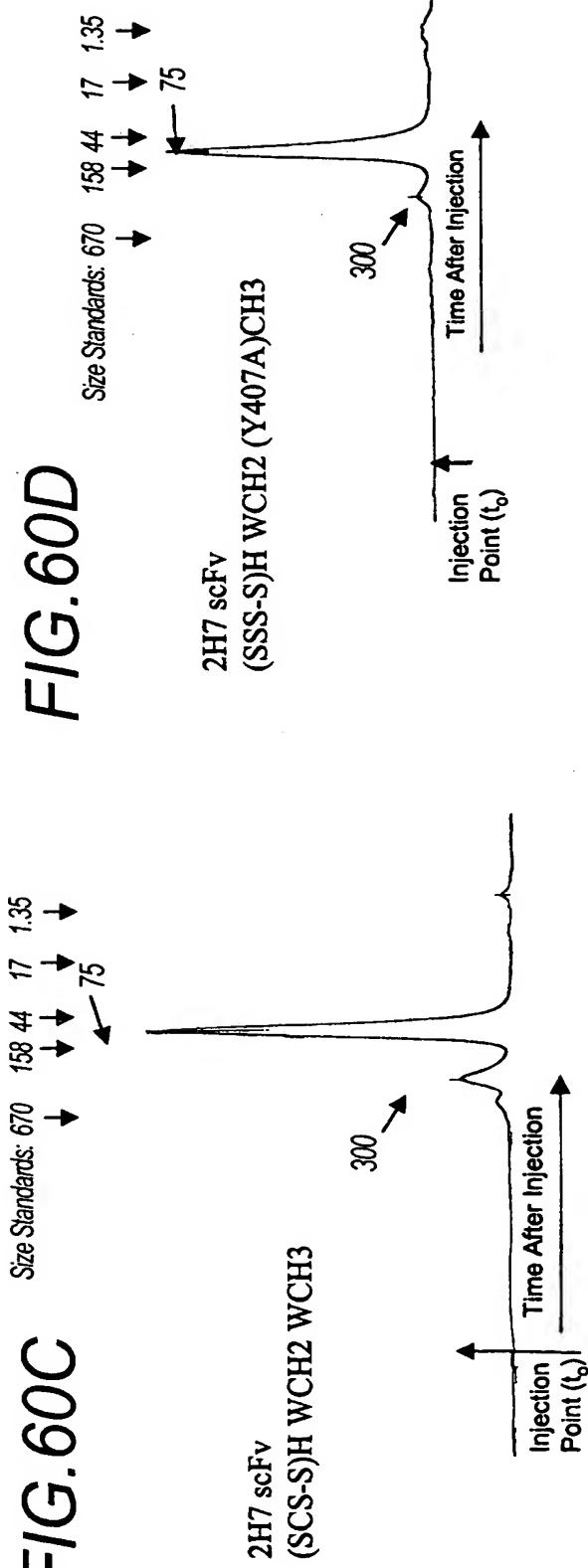


FIG. 60B

FIG. 60D

FIG. 61A

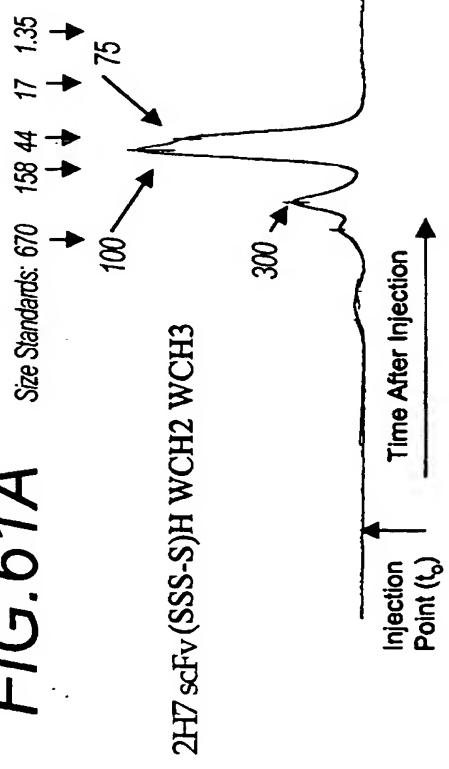


FIG. 61B

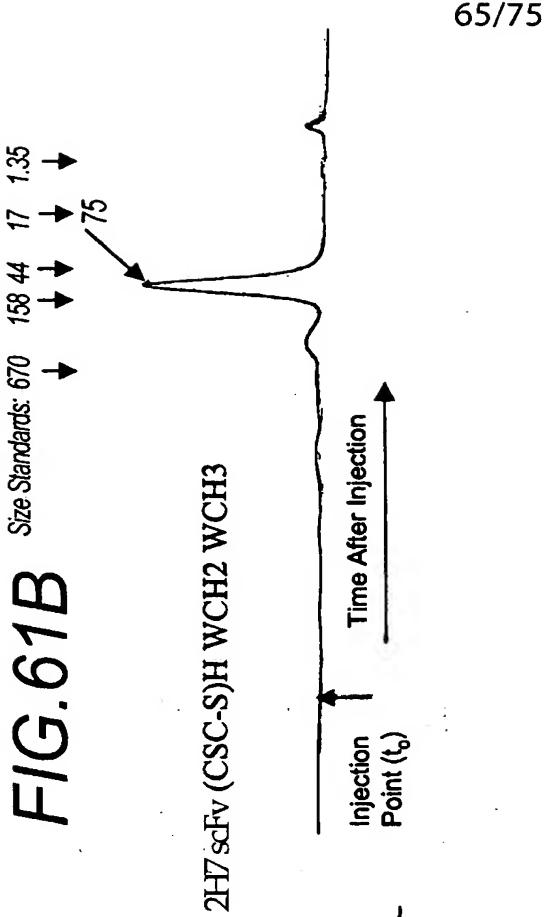


FIG. 61C

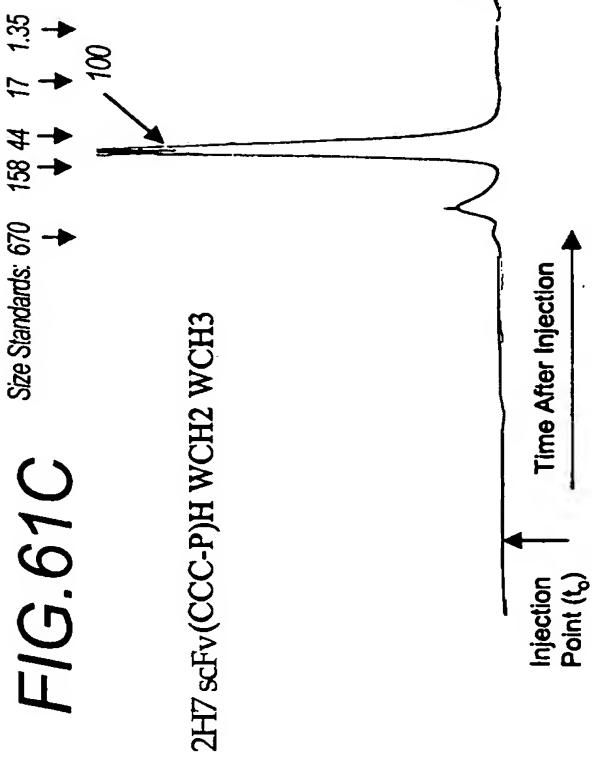
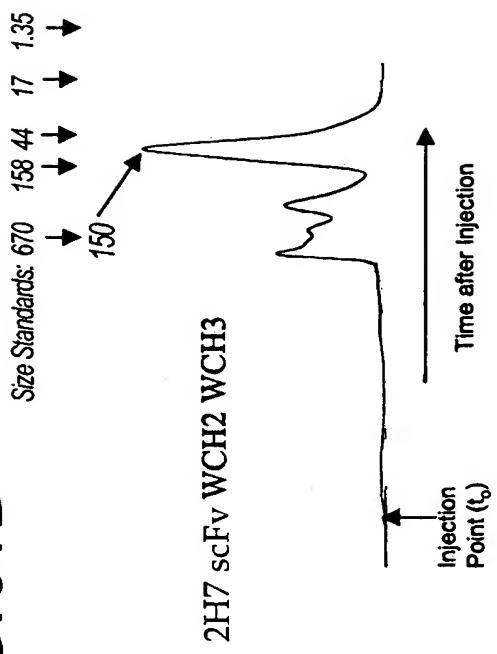


FIG. 61D



65/75

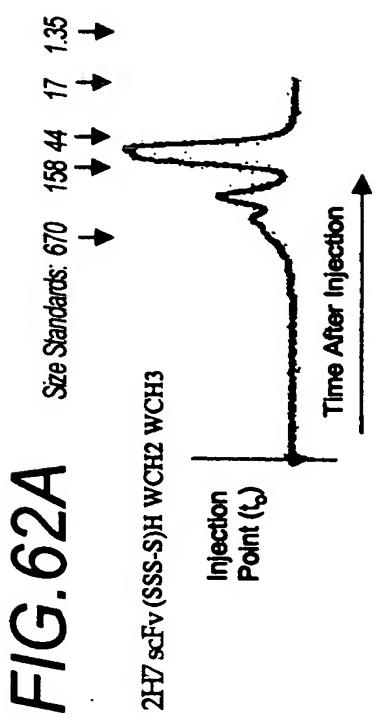
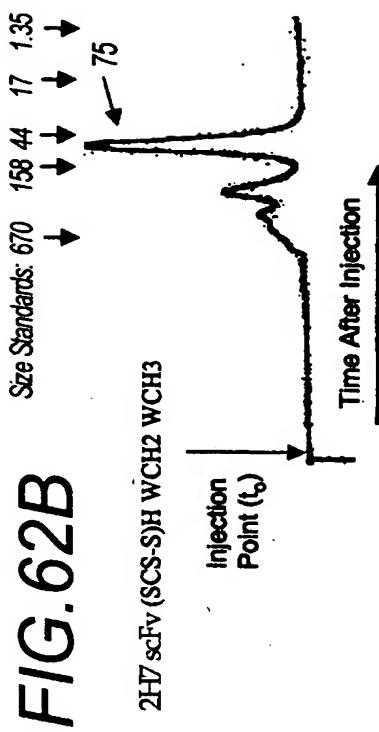


FIG. 62B



66/75

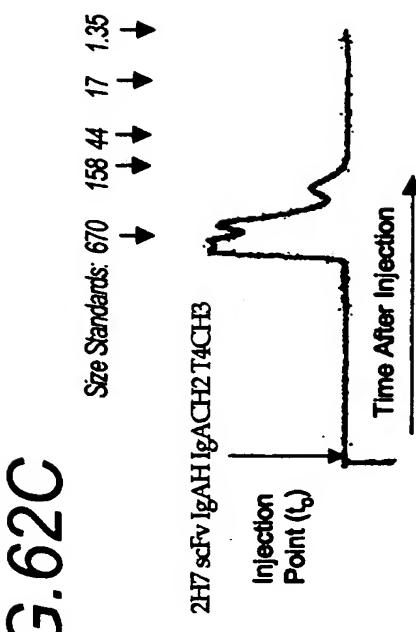


FIG. 62C

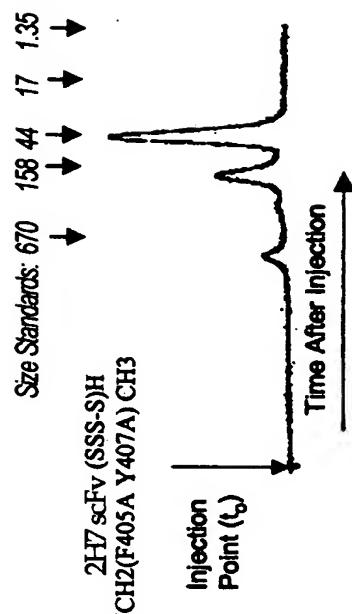


FIG. 62D

FIG. 63

Binding of Purified Proteins from COS Supernatants
to CD20 CHO cells:
Differential Effects of CH3 Mutations on Binding

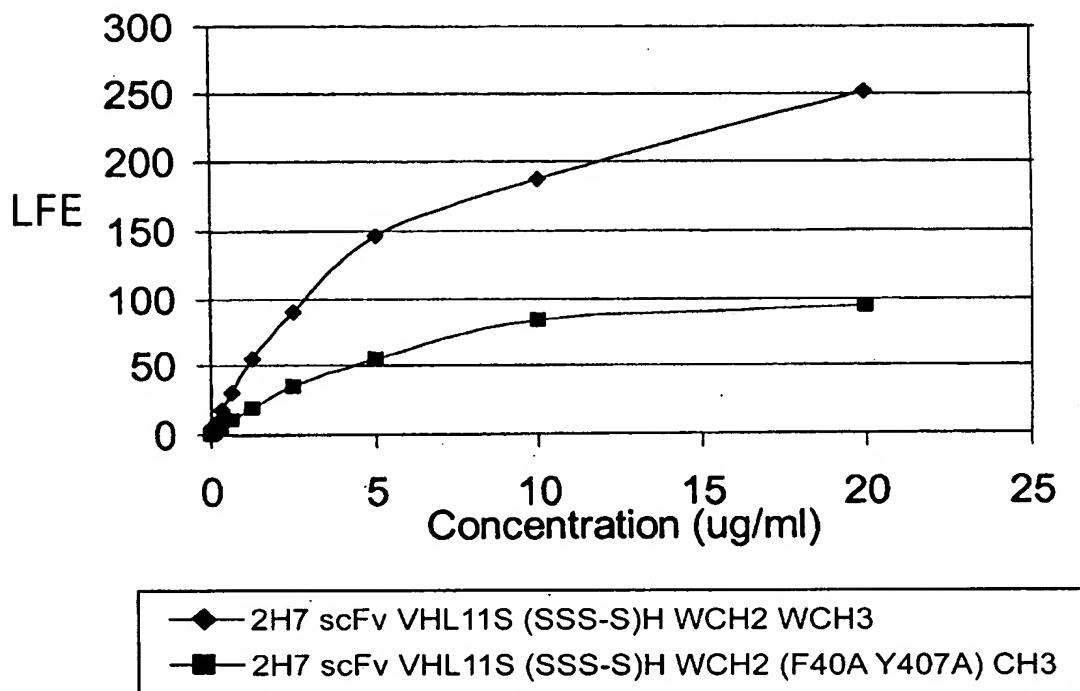


FIG.64

Binding of FITC conjugated 2H7 scFv VHL11S Proteins to CD20 CH0 Cells

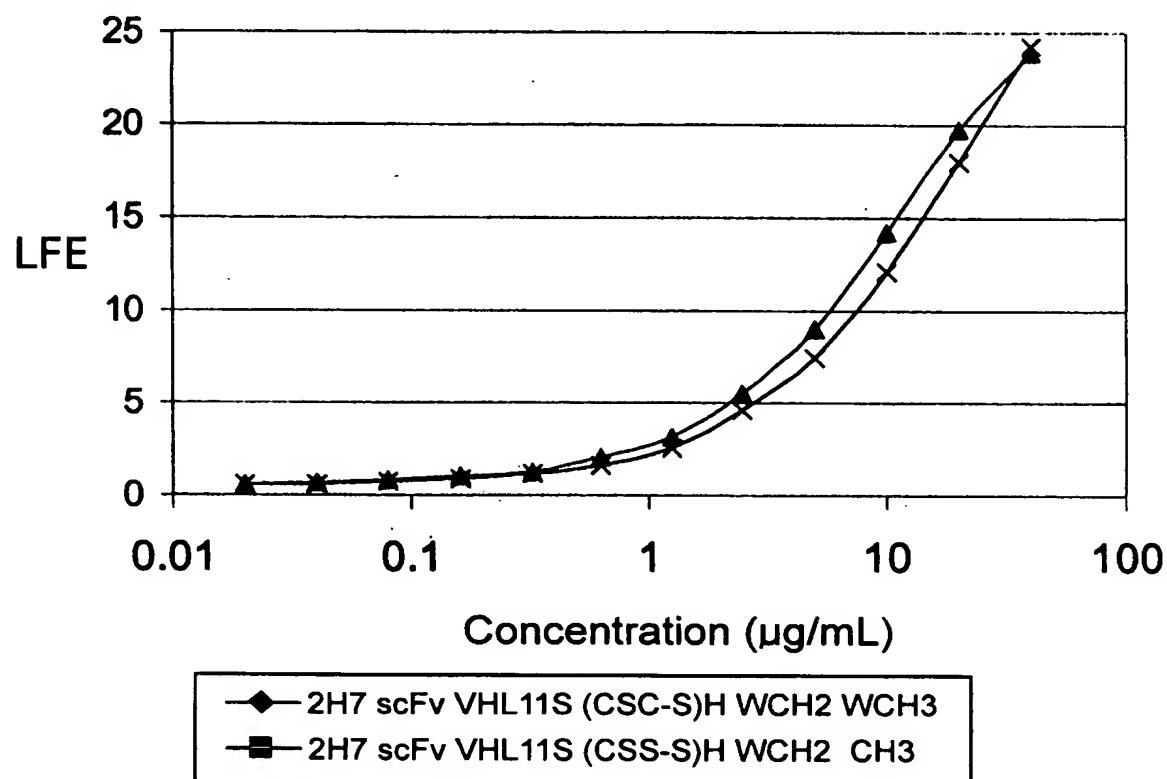


FIG. 65

Nonreducing SDS-PAGE on Protein A-Purified Lots
of 2H7 scFv VHL11S Constructs (10 ug/lane)

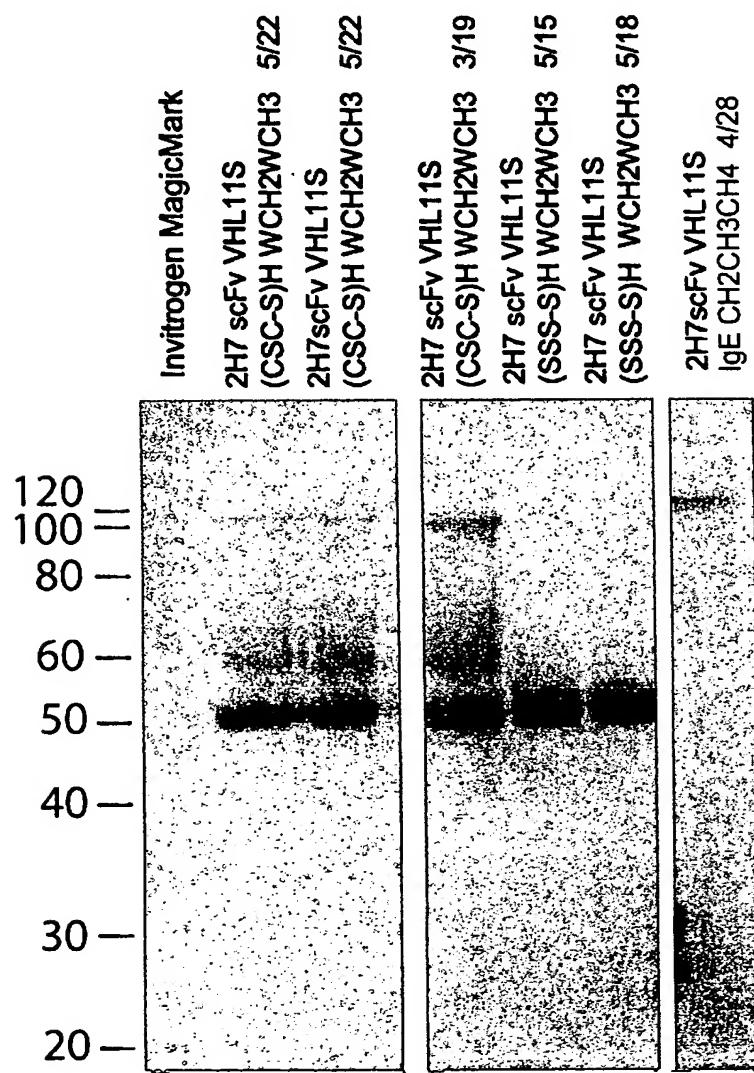


FIG. 66

Alterations in Human IgG Fc sequence
that differentially change effector function efficiency

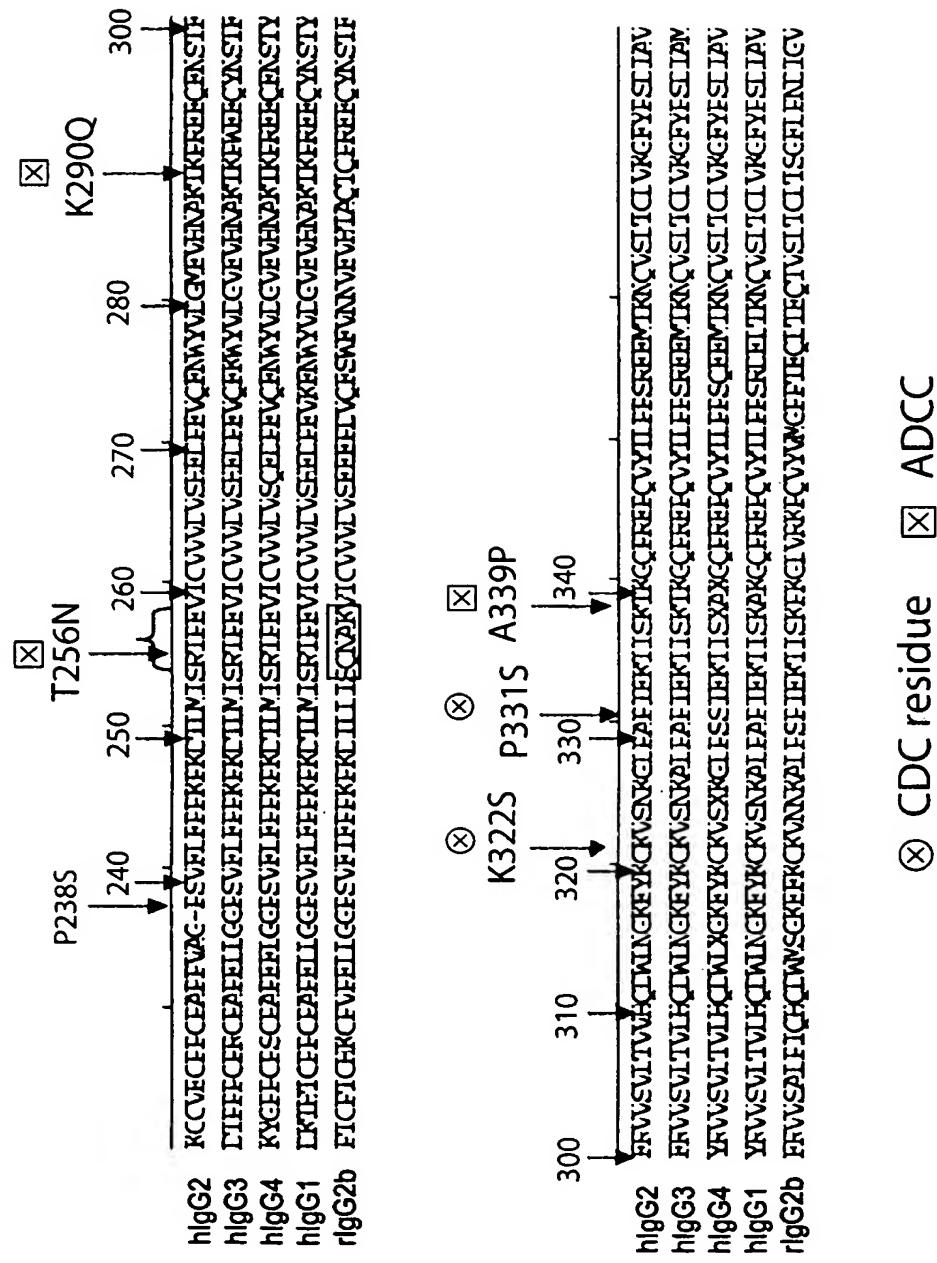


FIG. 67

ADCC Activity of 2H7 scFv VHL11S (CSC-S)H WCH2 WCH3 from CHO and Lec13-CHO transient transfections

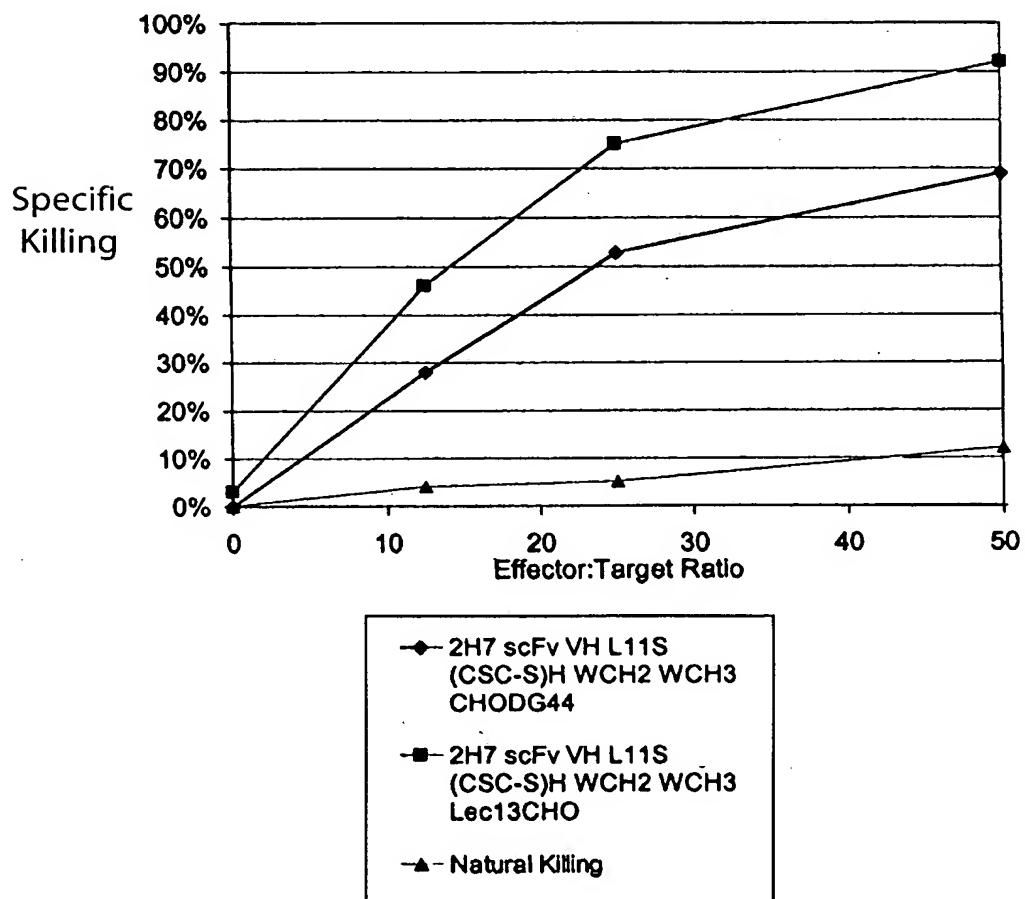


FIG. 68

CD16(ED)(SSS-S)H P238S CH2 WCH3 high and low affinity alleles expressed as soluble molecules

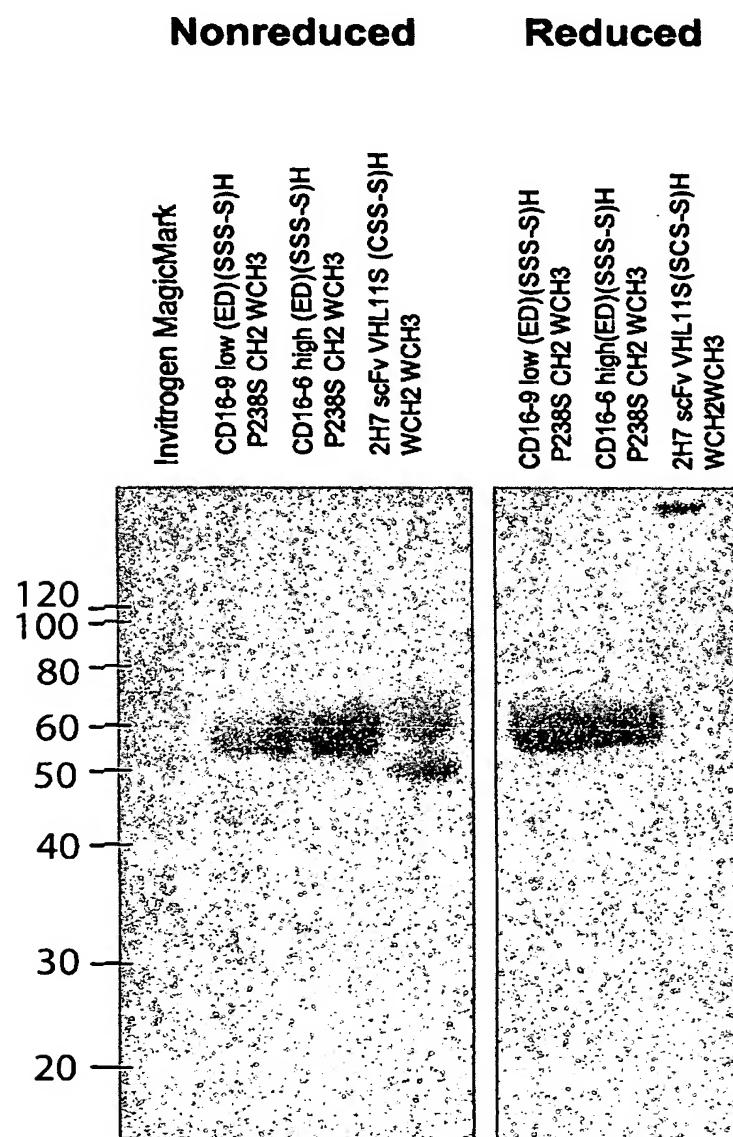


FIG. 69

**Binding of soluble CD16-FITC high and low affinity fusion proteins
to 2H7 scFv VHL11S (CSC-S)H WCH2WCH3 or
(SSS-S)H P238S CH2WCH3 on CD20CHO Targets**

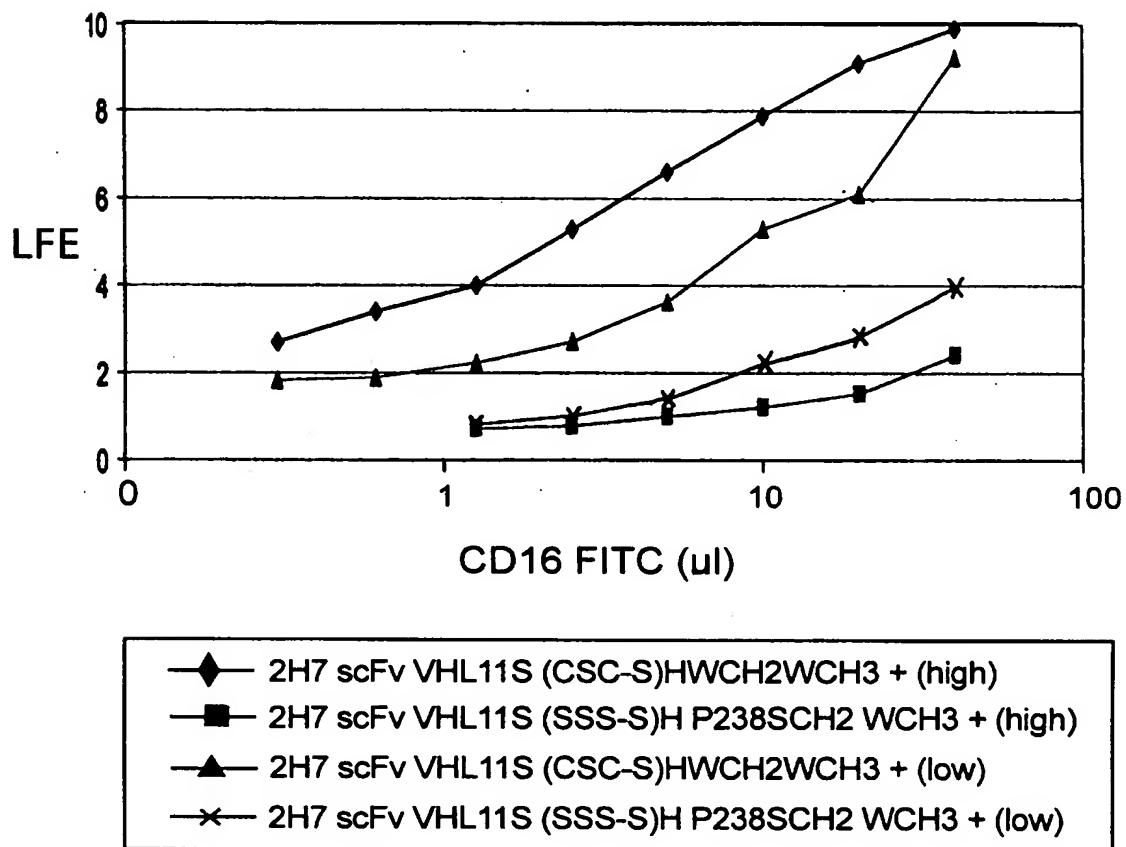
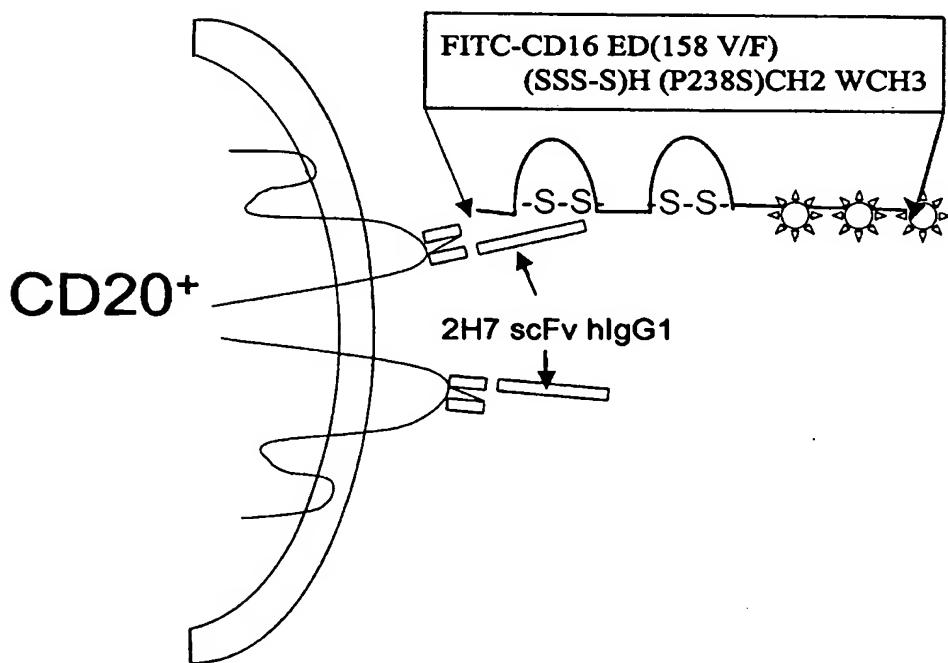


FIG. 70

Binding of FITC Labeled, Recombinant Human CD16(ED) extracellular domain -Ig Fusion Protein to CytoxB Derivatives on CD20 CHO Cells



Expression of surface displayed SMIPs links modified cDNAs with the altered fusion proteins

Mammalian Cell Transfected With
1.A single surface displayed scFvlg expression construct
OR
2.a library of such molecules

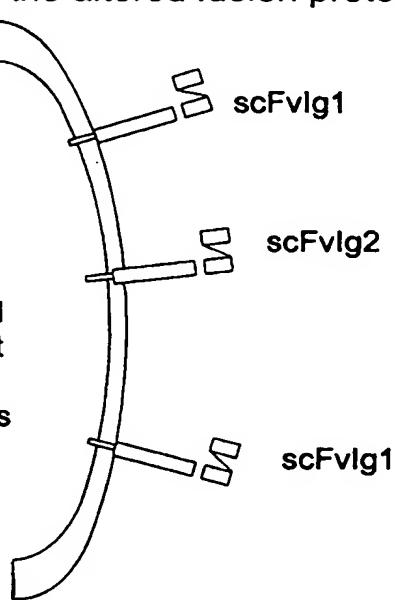


FIG. 71**CD37 mAbs and scFvIg Induce Apoptosis**

Bjab Staining	Annexin V Positive	
No scFvIg	17.5	
2H7 MH	27	
G28-1 MH	30.6	
G28-1 IgAH	28.9	
HD37 MH	29.1	
(2H7+G28-1)MH	41	
(2H7+HD37) MH	37.1	
(G28-1+HD37) MH	35.3	
		plus GAM
Ramos	AnnexinV Positive	AnnexinV positive
cells alone	3	3.3
2H7 Mab	1.4	3.1
G28-1 Mab	18.3	8.7
mAbs	HD37 Mab	3.7
	G28-5	3.9
	2H7+G28-1	32.3
	2H7+HD37	5
	2H7+G28-5	5.7
	HD37+G28-1	26.9
	HD37+G28-5	8.2
	G28-1+G28-5	39.5
		68.3